CURRENT FILING DATE:

2002-12-06

APPLICATION NUMBER: 60/336,695 FILING DATE: 2001-12-07 APPLICATION NUMBER: 10/226,294

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LENGTH: 177
TYPE: PRT
ORGANISM: human
US-10-202-062-40
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; ORGANISM: human
US-10-310-793-44
                                                                                                                                                                                                                                                                                                    Sequence 40, Application US/10202062
Publication No. US20040038349A1
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.,
TITLE OF INVENTION: Heteromultimeric TNF Ligand Family members
FILE REFERENCE: PF559
CURRENT APPLICATION NUMBER: US/10/202,062
CURRENT FILING DATE: 2002-07-25
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Best Local :
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Best Local :
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LENGTH: 177
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PRIOR FILING DATE: 2001-07-27
NUMBER OF cen
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                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 42
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FILING DATE: 2000-04-27
APPLICATION NUMBER: 60/180,908
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APPLICATION NUMBER: 60/314,381
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                                                            al Similarity
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                                                        100.0%; Score 951; DB 15; ilarity 100.0%; Pred. No. 2e-92; Conservative 0; Mismatches 0;
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Pred. No. 2e-92;
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US-09-345-790-2
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Best Local Similarity
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APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Human Endokine Alpha
NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/345,790
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
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20005-3934
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US20020099198A1
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hypothetical prote replication protei transcription acti interferon gamma p glycoprotein VP7 p glycoprotein VP7 p glycoprotein VP7 p neurotoxin type F hypothetical prote hypothetical prote probable glucan sy hypothetical prote penicillin-binding penicillin-binding probable purine nu hypothetical prote

ALIGNMENTS

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E.V.,

C;Decies: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Date: 13-Nov-1998 #sequence_revision 13-Nov-1998 #text_change 09-Jul-2004
C;Accession: A71621
R;Gardner, M.J. Tettelin, H.; Carucci, D.J.; Cummings, L.M.; Aravind, L.; Koonin, I.; Pertea, M.; Salzberg, S.; Zhou, L.; Sutton, G.G.; Clayton, R.; White, O.; Smith, Science 282, 1126-1132, 1998
A;Title: Chromosome 2 sequence of the human malaria parasite Plasmodium falciparum. A;Accession: A71600; MUID:99021743; PMID:9804551
A;Accession: A71621

A; Residues: 1-1188 'GAK' A; Cross-references: UNIPROT:096143; GB:AE001380; GB:A; Cross-references: UNIPROT:096143; GB:AE001380; GB:A; Cross-references: UNIPROT:096143; GB:AE001380; GB:A; Cross-references: UNIPROT:096143; GB:AE001380; GB:A; Cross-references: UNIPROT:01307 C; Genetics: A; Gene: PFB0205c Qy		N N N N N	G C C F H B	Дb	γQ	Дb	γQ	상 성		≯C;	P P P P	PP
Length 1188; Length 1188; 5; Indels 40; Gaps 9; 6(VNKVSDWKLEIL 90	A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-665 <wil> A;Residues: 1-665 <wil></wil></wil>	R;Percy, C. submitted to the BMBL Data Library, July 1996 submitted to the BMBL Data Library, July 1996 A;Reference number: Z19056 A;Accession: T18979 A;Status: preliminary; translated from GB/EMBL/DDBJ	RESULT 2 T18979 hypothetical protein C06B3.2 - Caenorhabditis elegans c;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004 C;Accession: T18979	112 INNTY			91	41 FSWLIFIFLOLETAKEDCMAKFG-PLPSKWQMASSEPPCVNKVSDWKLEIL	8.6%; Score 82; DB 2; Length 1188; pal Similarity 30.4%; Pred. No. 19; 38; Conservative 11; Mismatches 36; Indels 40; Gaps	A;Gene: PFB0205C	A;MOLECULE Cype: DNA A;Residues: 1-1188 <gar> A;Cross-references: UNIPROT:096143; GB:AE001380; GB:AE001362; NID:g3845120; PIDN:AAC718 A:Experimental source: clone 3D7</gar>	A;Scatus: preliminary; nucleic acid sequence not shown; translation not shown

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A; Introns: 23/1; 202/2;
A; Note: F2K13_50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Experimental source: cultivar Columbia; BAC clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: Z25394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Sato, S.; Nakamura, Y.; Kaneko, submitted to the Protein Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor protein kinase-like protein - Arabidopsis thaliana
N;Alternate names: protein F2K13 50
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 16-Aug-2004
C;Accession: T51545
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession: T51545
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A;Introns: 29/3; 91/1; 139/3; 205/2; 240/1; 296/1; 337/3; 390/1; 419/3; 483/2; 556/2; C;Superfamily: Caenorhabditis elegans hypothetical protein F09B9.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B
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A;Gene: CESP:C06B3.2
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                                                                                                                                                                                                                                                                                                                                                                                                   Superfamily: protein kinase homology
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                                                                                         175
                                                                                                                        160
                                                                                                                                                    120 LYLGPNKWVRVDLEGK-
                                                                                                                                                                                                       63 QKNLEAVHIKEYLFLRYFPDGLRNCYTLDVLQNRRYMIKAVFVYGNY---DGYNDYPSFD
                                                                                                                                                                                                                                                                                                     43 WIIFIFIQLETAKEPCMAKFGPLPSKWQMASSEPPCVNKV-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            151
                                                                                                                                                                                                                                                                    7 YÉFFÍFAIIHYVQ----AQQGFISLDCGLPSNEÞPYIEFVTGLVFSSDADHIPSGISGRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    120 NKDMIQTL---TN----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56 SSIEQFTAASSDCLFFKNCTASILNILKDNIYAIQQYDAFAKIESAGLLEVSLVFDGSYQ 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1-851 <SAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27 WIFCSIVMLLFLCSFSWLIFIFLQLETAKBFCMAKFGPLFSKWQMASSBFPCVNKVSDW-
                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44;
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                                                                                                                     KNNTY 164
                                                                                                                                                                           VRLYKNKDMIQTLINKSKIQNVGGTYE--LHV-----
                                                                                         RNDTY 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOLSALPLTACATECSORDVPKDSAFWG 203
                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                 8.4%; 5cc-
y 22.2%; Pred
ye 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Kaneko, T.; Kato, T.; Asamizu,
Sequence Database, August 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    --IFNSEHQVLKNNTYWG 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.4%; Score 79.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -----KLEILQNGLYLI--YGQVA--PNANYNDVAPFEVRLYK 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24; Mismatches
                                                                                                                                                                                                                                                                                                                                                       Score 79.5;
Pred. No. 22;
                                                                                                                                          VNGSVEEIIHIPSSNSLQICLVKTGNSLPFISALELRLL 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 16;
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                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                   --WKLEILQNGLYLI----YGQVAPNANYNDVAPFE 114
   Mycoplasma
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                                                                                                                                                                                                                                                                                                                                      44;
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   pneumoniae
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                                                                                                                                                                         GDTIDLIFNSEHQVL
 (strain
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ATCC
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29342)
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C,Superfamily: oligopeptide permease protein oppB
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probable peptide ABC transporter, permease protein VCA0589 [imported] - Vibrio cholerae C;Species: Vibrio cholerae C;Species: Paug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004 C;Accession: D82442
                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q9KM00; GB:AE004389; GB:AE003853; NID:g9657989; PIDN:AAF9649]
                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-352 <HEI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae A;Reference number: A82035; MUID:20406833; PMID:10952301
A;Accession: D82442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J. chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, I., R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
                                                                                                                                                                                                                                                                                                                                                       C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Status: preliminary
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C;Species: Mycoplasma pneumoniae
A;Variety: ATCC 20342
C;Date: 26-Feb-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S73405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Gene: uhpT
A;Genetic code: SGC3
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A;Note: the mucleotide sequence was submitted to the EMBL Data Library, November 1996
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-564 < HIM>
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A;Status: preliminary; nucleic acid sequence not shown; translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Himmelreich, R.; Hilbert, H.; Plagens, H.; Pirkl, E.; Li, B.C.; Herrmann, R. Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma pneumoniae
A;Reference number: $73327; MUID:97105885, PMID:8948633
                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                 Gene: VCA0589
                                                                                                                                                                                                                      Query Match
Best Local :
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Best Local Similarity
Matches 33; Conserv
                                                                                                                                                                                                                 Local Similarity
                                                         67
                                                                                                 14
                                                                                                                              10 PLSH---SRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLP 66
           61 GAWYM--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           117 LYKNKDMIQTLTNKSKIQNVGGTY 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   288 LGNQPSAGDILKRKATWKMIGMYGICLVVLVNPLTGGWWNILQAVSPASSFN---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       238 QSVQQAGTKYWVFILTALILIGFGILCVPAW-----FEKQMDPFMPQ----KQTKEQMQ 287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
                                      SKWOWASSEPPCVNKVSDWKLEILONGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMI 124
                                                                                   PLTEARWARFKANRRGFWSLWIF----LLLFVVS-----LFAELIANDKPLLIQY---D 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ---VKDGVKTL---KPLEGAGGYF 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSEPPCVNKV----SDWK-----LEILQN----GLYLIYGQVAPNANYNDVAPFEVR 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QGAQRSSWKLWLFCSIVMLL----FLCSFSWLIFIFLQLETAKEPCMAKFGPLPSKWQMA 72
                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                            8.1%; Score 77.5;
26.3%; Pred. No. 12;
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22.9%; Pred. No.
tive 22; Mismatc
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19;
GEFDTEADYTD -- PYVVSLIEEKGQI 101
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K;Messenguy, F.; Dubois, E.; Descamps, F.
Eur. J. Biochem. 157, 77-81, 1986
A;Title: Nucleotide sequence of the ARGRII regulatory gene A;Reference number: A25064; MUID:86220196; PMID:3709534
A;Accession: A25064
                                                                                                                                                                                                                                                    glycoprotein VP7 precursor - human rotavirus A (serotype 2 strain DS1)

N,Alternate names: outer capsid protein VP7

C;Species: human rotavirus A
C;Date: 30-Jun-1989 #sequence_revision 30-Jun-1989 #text_change 09-Jul-2004
C;Accession: E27620
R;Green, K.Y.; Midthun, K.; Gorziglia, M.; Hoshino, Y.; Kapikian, A.Z.; Chanock, R.M.;
Virology 161, 153-159, 1987
A;Title: Comparison of the amino acid sequences of the major neutralization protein of A;Accession: E27620
A;Accession: E27620
A;Accession: E27620
A;Molecule type: genomic RNA
A;Residues: 1-326 <GRE>
A;Genetics:
UNIPROT:P11850
C;Genetics:
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A;Map position: 13L
C;Superfamily: GAL4 zinc binuclear cluster homology
C;Keywords: DNA binding; nucleus
F;16-53/Domain: GAL4 zinc binuclear cluster homology
                                                                                                                                                        A;Map position: segment 9
C;Superfamily: rotavirus glycoprotein VP7
C;Keywords: coat protein; glycoprotein; transmembrane protein
F;1-23/Domain: signal sequence #status predicted <SIG>
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VGXRDS
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A;Residues: 1-3,'F',5-128,130-131,'L',132-282,'V',284-344,'V',346-365,'Q',367-548,'A',5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    regulatory protein ARGR2 - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein YML099c
C;Species: Saccharomyces cerevisiae
C;Date: 02-Dec-1994 #sequence_revision 10-Feb-1995 #text_change
C;Accession: S49677; A25064
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A; Accession: S49627
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:P05085; EMBL:Z46660; NID:g575702; PIDN:CAA86638.1; R;Messenguy, F.; Dubois, E.; Descamps, F.
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                                                                                      11-23/Domain: signal sequence #status predicted <SIG>;24-326/Product: glycoprotein VP7 #status predicted <GPV>;32-48/Region: hydrophobic #status predicted (GPV);69,146,238/Binding site: carbohydrate (Asn) (covalent) #
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;Residues: 1-880 <GEN>
       Matches
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Best Local
                            Local
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    42;
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                            Similarity
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  Conservative
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                          22.8%;
                                             8.1%;
    29;
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                        Score 77; DB
Pred. No. 12;
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    59;
                                             Length 326
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R;Green, K.Y.; Midthun, K.; Gorziglia, M.; Hoshino, Y.; Virology 161, 153-159, 1987
A;Title: Comparison of the amino acid sequences of the the A;Reference number: A27620; MUID:88044489; PMID:2823458
                                                                                   C;Species: human rotavirus A
C;Date: 30-Jun-1989 #sequence revision 30-Jun-1989
C;Accession: G27620; A93520; A04134
                                                                                                                                                                                       RESULT
VGXRHU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           F;24-326/Product: glycoprotein VP7 #status pro
F;32-48/Region: hydrophobic #status predicted
F;69,146,238/Binding site: carbohydrate (Asn)
                                                                                                                                            glycoprotein VP7 precursor - human rotavirus N;Alternate names: outer capsid protein VP7
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F;1-23/Domain: signal sequence #status predicted <SIG>
F;24-226/Product: glycoprotein VP7 #status predicted <GPV>
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A;Title: Comparison of the amino acid sequences of the major neutralization protein of A;Reference number: A27620; MUID:88044489; PMID:2823458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: human rotavirus A
C;Date: 30-Jun-1989 #sequence_revision
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A; Residues: 1-326 < GRE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: UNIPROT: P11851
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                                                                                                                                                                                                                                                                                                                                                      130 NPQLYCDYNVV
                                                                                                                                                                                                                                                                                                                                                                                               116 --RLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LWLFCSIVMLLFLC----SFSWLIFIFLQLETAKEPCMAK-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CV-----NKVSDWKLEILQNGLYLIYGQVAPNA---NYNDVAPFEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29;
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                                                                                                                                                                                                                                                                                                                                                      LMRYDNTSELDASELADLILNE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 77; DI
Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-Jun-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (covalent) #status
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                                                                                                      #text_change
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                                                           Y.; Kapikian,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 326
                         major neutralization
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                                                                                                                                                                                                                                                                                                                                                      W----LCNP
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                                                               A.Z.; Chanock,
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A; Accession: G27620
A; Molecula ---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;McGraw, R.A.; Coffee, B.W.; Otto, C.M.; Drews, R.T.; Rawlings, Nucleic Acids Res. 18, 5563, 1990
A;Title: Gene sequence of feline tumor necrosis factor alpha. A;Reference number: S11688; MUID:91016860; PMID:2216740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumor necrosis factor alpha precursor - cat
C;Species: Felis silvestris catus (domestic cat)
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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F;24-326/Product: glycoprotein VP7 #status predicted <GPV>
F;32-48/Region: hydrophobic #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Experimental source: serotype C; Genetics:
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A; Residues: 1-326 <GRE>
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                                                                                                                                                                                                                                                                                                                                                                                                                             A; Introns: 62/3; 78/1; 94/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: DNA
A;Residues: 1-233 <MCG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: S11688
                                                                                                                                                                                                                                                                            ;Superfamily: tumor necrosis factor; ;Keywords: glycoprotein; lipoprotein; myristylation; transmembrane;18,20/Binding site: myristate (Lys) (covalent) #status predicted;81/Binding site: carbohydrate (Ser) (covalent) #status predicted;81/Binding site: carbohydrate (Ser)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Molecule type: DNA
                                                                                                                                                                                                                                               ;145-177/Disulfide bonds: #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: UNIPROT:P19101; EMBL:X54000; NID:g1084; PIDN:CAA37948.1; PID:g29577
                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42;
                                       7
                                                                                                         Similarity 50; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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ENMPLSHSRTQGAQRSSWKLMLFCSIVMLLFLCSFSWLIF-----IFLQLETAKEPCMAK 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MDIS 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QFIS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   --RLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASSEPP-----NYNDVAPFEV 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STŚGEPFLISTLCLYYPAEAKNEIŚDDEWENTLSQLFLTKGWPIGSVYFKDYNDINTFSV 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LTILISIILLNYILKTITNTMDYIIFRFLLLIALISPFVRTQNYGMYLPITGSLDAVYTN
                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                         8.0%;
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                                                                                                     31;
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                                                                                                                                             Score 76.5;
Pred. No. 9.
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                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                      9.2;
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                                                                                                         54;
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                                                                                                                                                                      2;
                                                                                                                                                                          Length 233;
                                                                                                         Indels 113;
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                                                                                                            12;
          A; Cross-references:
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A86736
C; Species: Lactococcus lactis subsp. lactis (strain C; Species: Lactococcus lactis subsp. lactis
C; Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 09-Jul-2004
C; Accession: A86736
                                                                                                                                                                                                                                               hypothetical protein BB0205 - Lyme disease spirochete C;Species: Borrelia burgdorferi (Lyme disease spirochete) C;Decies: Borrelia burgdorferi (Lyme disease spirochete) C;Date: 13-Feb-1998 #sequence_revision 13-Feb-1998 #text_change 09-Jul-2004 C;Accession: E70125
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E70125
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A;Accession: A86736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R; Bolotin, A.; Wincker, P.; Mauger, Genome Res. 11, 731-753, 2001
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A; Molecule type: DNA
A; Residues: 1-525 < KLE>
                                                                       A; Reference number: A; Accession: E70125
                                                                                         A;Authors: Smith, H.O.; Venter, J.C.
A;Title: Genomic sequence of a Lyme disease spirochaete,
A;Reference number: A70100; MUID:98065943; PMID:9403685
                                                                                                                                                                                 son, D.; Peterson, J.; Bowman, C.; Garland,
                                                                                                                                                                                                                      R; Fraser,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
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                                              A;Status: preliminary; nucleic acid sequence
                                                                                                                                                          Nature 390,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                     C.M.; Casjens, S.; Huang, W.M.; Sutton, G.G.; Clayton, R.; Lathigra, R.; Peterson, J.; Kerlavage, A.R.; Quackenbush, J.; Salzberg, S.; Hanson, M.; C.; Garland, S.; Fujii, C.; Cotton, M.D.; Horst, K.; Roberts, K.; Hatch, J., 580-586, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        163
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                                                                                                                                                                                                                                                                                                                                                                                                                                LDDVTFKPSGNSLSIKKMTGNTKIIVPEDVGVSLDLTTNAGLVKIFNEAAQINAGN 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KFGPLPSKWQMASSBPPCVNKVSDWKLEILQNGL-YLIYGQVAP-----NANYNDVAP 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    I--KSPCQRETPEGAEAKPWYEPIYLGGVFQLEKGDRLSTEINLPAYLDFAESGQV----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FGVIGPQREELPHGLQLINPLPQTLRSSSRTPSDKPVAHVVANPEAEGQLQRLSRRANAL 112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UNIPROT:Q9CH50; GB:AE005176; PID:g12723820; PIDN:AAK04987.1; GSPDB:Gb
se: strain IL1403
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Pred. No.
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                                                                                                                     Borrelia burgdorferi
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                                                                                                                                                                                     White,
Vugt,
B.
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UNIPROT:051223; GB:AE001131; GB:AE000783; NID:g2688098; PIDN:AAC66601

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RESULT 14

ISOS11

MHC class I protein, alpha 1, alpha 2, alpha 3 and transmemu.
C.Species: Latimeria chalumnae (coelacanth)
C.Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_cha:
C.Accession: ISOS11
R.Betz, U.A.K.; Mayer, W.E.; Klein, J.
R.Betz, U.A.K.; Mayer, W.E.; Mein, J.
Proc. Natl. Acad. Sci. U.S.A. 91, 11065-11069, 1994
A;Title: Major histocompatibility complex class I genes of A:Reference number: ISOS10; MUID:95062206; PMID:7972010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Gene: dal2
C;Superfamily
C;Keywords: DI
F;2-57/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MADS-box protein (C;Species: Picea (C;Date: 14-Jul-19) (C;Accession: T148)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: DNA
A;Residues: 92-156 <TA2>
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A; Residues: 1-222 < TAN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Superfamily: transcription factor squa; serum response factor DNA-binding Keywords: DNA binding; transcription regulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANDS-box protein dal2 - Norway spruce Species: Picea abies (Norway spruce) Date: 14-Jul-1995 #sequence_revision Accession: T14847; S51934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                         MIQILINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNN
                                                                                                                                                                                                                                                                                                            ESNSQYWQQEAGK-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A binding; transcription regulation serum response factor DNA-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source: strain B31
Borrelia burgdorferi hypothetical protein BB0205
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                                                                                                                                                                                                                                                                                                                                          -WQMASSEPPCVNKVSDWKLEILONGLYLIYGQVAPNANYNDVAPFEVRLYKNKD
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                                                                                                                                                                                                                                                                                                          -LRQQIEILQNANRHLMGDGLTALNIKELKQLEVRLEKGIG
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Pred. No. 11
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A;Molecule type: DNA
A;Residues: 1-686 <WIL>
A;Cross-references: UNIPROT:Q19467;
             Search completed: November
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A; Introns: 9/1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                submitted to the EMBL Data A; Reference number: Z19343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Kershaw,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein F14F3.2 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #t
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C; Superfamily: class I histocompatibility antigen;
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A;Status: preliminary; translated
A;Molecule type: DNA
A;Residues: 1-314 <BET>
A;Cross-references: UNIPROT:Q31417
C;Genetics:
                                                                                                                                                                                                                                                                                                                                               A; Experimental source:
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Tocal Similarity
time
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43
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                                                                                                                                        MASSEPP-CVNKVSDWKLEILQNGLYLIYGQVA-----PNANYN------ 108
                                                                                                                                                                     MSAARNQRRQK------359
                                                       DELLEMKERLSSTETMVVTLTK---
                                                                               -DVAPFEVRLYKNKDMIQTLINKSKIONVGGTYELHVGDTIDLIFNSEHQVLKNNTY 164
                                                                                                            -PVSEPPKNMRKSNDTPLRIMKKSEYADYDEVAGISCSPISGIVPNSQANTLDESSSRVW 418
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49/1; 76/2; 100/2; 151/2; 212/3; 314/3; 380/2; 447/2;
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34.2%;
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Pred. No. 16;
9; Mismatches
                                                                                                                                                                                                                                            Score 75.5;
Pred. No. 41;
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                                                    -QNEELTKLVHSLQAQNINLNSELITFRDDLY
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AAP60547 ADE36761 AAW10344 AAO21502

Aao21502

Aag62280 Abb83970 Abp35593 Abb62664 Aag43532 Aag43532 Aag43531 Aag43531 Aag60547 Ade36761

Propionib
Novel pro
Streptoco
Streptoco
Human gen
Human U2
Fungal ZB
Fungal

ADC94332 AAB18183 AAU57900 ABM54419 ABM65418 ABM93512 ADF2593 ADF2593 AAG62280 ABF2593 AAG62280 ABB2593 ABB62664 ABB63664 ABA643533 AAG43533 AAG43533

Aau57900 Abm54419

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Abm64618 Abb93512 Abp35594 Ade09068 Abp25933

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Minimum
Maximum
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                            score greater than and is derived by
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BB
                                                                                                                                                                                                                                                                                                                  seq
                                                             No.
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                            is the number of results predicted by chance to have ater than or equal to the score of the result being prived by analysis of the total score distribution.
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1: geneseqp1980s:*
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                                                                                                                                                                                                                                                                                      Minimum Match
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geneseqp2004s:*
                                                                                                                                                                                                                                                                                                                                                                                                 seqs,
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SUMMARIES
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412.306 Million cell updates/sec
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(without alignments)
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	ALIGNMENTS
RESULT	ሆነ 1
AAY15817	
ID	AAY15817 standard; protein; 177 AA.
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AC	AAY15817;
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ď	28-JUL-1999 (first entry)
××	
DE	Amino acid sequence of human DNA19355 polypeptide.
×	
KW	DNA19355; tumour necrosis factor homologue; receptor GITR; immunogen;
K¥	apoptosis; mammalian cancer cell;
XX	tumour necrosis factor (INF)-alpha secretion; primary I- cell;
ž ž	proinflammatory response.
20 Y	Homo sapiens.
7	W0000E024 - 31
XX PN	WO9925834-A1.
PD	27-MAY-1999.
3	
YY PF	18-NOV-1998; 98WO-US024621.
3 }	
Яď	18-NOV-1997; 970S-0065635P.
PR	12-DEC-1997; 97US-0069661P.
PA XX	(GETH) GENENTECH INC.
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DR XX	WPT: 1999-338009/28.
DR	N-PSDB; AAX59744.
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Result

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Score

Match Query

Length

BB

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AAY15817 AAY06646 AAB47056 AAB20109

100.0 100.0

AAB53065 AAB47287 AAB50979 AAU81953 ABR42323

Aab47287 Aab47056 Aab20109 Aay15817 Aay06646 Description

Human a PRO175

Human imm PR0175.

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Human Human

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Human

Claim 10; Fig 1; 86pp; English.

New DNA19355 polypeptide as tumour necrosis factor homolog

Human Human

ABP60544 ADC35222 ABW02285

Aab50979
Aau81953
Abr42323
Abr42323
Abp60544
Adc35222
Abw02285
Aaw37847
Aab08786
Aaw373061
Aab211464
Adc011464

AAW37847 AAB08785 AAY5307 AAY5307 ADA11464 ADC01858 ABC17426 AAM70748 AAM38998 AAE14752 AAM14753 ADA17815 AAM41041

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The present sequence represents a polypeptide designated DNA19355. The polypeptide is a tumour necrosis factor homologue. The DNA19355 polyprocleotide sequence can be used to derive hybridisation probes for e.g. isolating similar sequences, gene mapping, genetic analysis, etc. Nucleic acids which encode DNA19355 can also be used to generate transgenic or knockout animals, which are useful in the development and screening of therapeutically useful reagents. The DNA19355 polypeptides may be used in diagnostic assays to detect the presence of the receptor GITR in mammalian tissues. The polypeptides can also be used as immunogens to raise antibodies. The polypeptides may also be used to

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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    뫄
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Best Local :
                                                                     Tumor necrosis factor receptor homologue - useful for, e.g. modulating apoptosis and N\bar{r}-KB activation and proinflammatory or autoimmune
                                                                                                                                                   WPI; 1999-494296/41.
N-PSDB; AAX87726.
                                                                                                                                                                                                                         Ashkenazi AJ, Goddard A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          induce apoptosis in mammalian cancer cells. DNA19355 polypeptides stimulate secretion of tumour necrosis factor (TNF)-alpha in primary cells, and so can be used to stimulate a proinflammatory response in mammalian cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-AUG-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRO364 ligand; tumour necrosis factor receptor; human; apoptosis; inflammation; antiinflammatory; NF-KB activation; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                     09-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                    09-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9940196-A1
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                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Domain
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nilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "N-glycosylation"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "transmembrane domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note= "cytoplasmic domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "N-glycosylation"
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Pred. No. 4.7e-97;
Mismatches 0;
                                                                                                                                                                                                                                                SA,
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Example 2; Fig 5A; 104pp; English

12-JUL-1999; 11-JUL-2000;

99US-0143304P

2000WO-US018867.

(GETH) GENENTECH INC

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                                                                                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                age-related macular degeneration; antibody; periodontal disease; vascular-related drug targeting; atherosclerosis; hypertension; inflammatory vasculitides; Reynaud's disease; aneurysm; arterial restenosis; thrombophlebitis; tumor angiogenesis; lung; fibrosis; neuropathy; rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRO364; human; glucocorticoid-induced tumor necrosis factor; recept hGITR; ligand; hGITRL; PRO175; tumour necrosis factor receptor; TNI human umbilical vein endothelial cell; HUVEC; cardiac hypertrophy; myocardial infarction; PGF_2alpha; trauma; cancer; angiogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 177
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                                                                                                                                                                                                                                                                                                                                                                                                           Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                           /note=
129
                                                                                                                                                                                                                                                                                                                                                                                                                                                              52. .177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                               /note= "N-glycosylated"
161
                                                                                                                                                                                                                                                                                                                 'note= "N-glycosylated"
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100.0%; I
ative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Transmembrane region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Cytoplasmic region"
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Pred. No. 4.7e-97;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             construction and expressed sequence tag (EST) database as having homology considered from an expressed sequence tag (EST) database as having homology converges of the tumour necrosis factor receptor (TNFR) family of color polypeptides. The PRO175 cDNA sequence was isolated from a library of cDNA fragments derived from human umbilical vein endothelial cells (HUVEC). Administering an effective amount of PRO364 or PRO175 or their antagonists is useful for treating cardiac hypertrophy (which is contained by myocardial infarction and characterized by the presence of initiated by myocardial infarction and characterized by the presence of macular degeneration in a human. Administering a therapeutically contained the firm a tumor or a natibody that binds PRO364 or PRO175 is useful for inhibiting angiogenesis induced by PRO364 or PRO175 in a human suffering from a tumor or a retinal disorder. PRO364 or PRO175, or their contained the firm and treatment or prevention of atherosclerosis, hypertension, inflammatory vasculatries, tumor angiogenesis, aneurysms, are useful for vascular-related drug targeting or as therapeutic targets for the treatment or prevention of atherosclerosis, hypertension, inflammatory vasculations, Reynaud's disease, aneurysms, and treatment of lung or liver fibrosis, periodontal diseases, attraction of bone-forming cells, central and peripheral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRO175 may be used in a mixture with a cardiovascular, endotherial, angiogenic or angiostatic agent for the treatment of a cardiovascular, angiogenic or angiostatic disorder. The PRO364 sequence was endothelial, angiogenic or angiostatic disorder. The PRO364 sequence was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents PRO175 polypeptide, which is a human gluco-corticoid-induced tumor necrosis falypertide (hGITRL). The cortrasponding receptor (hGITR), pRO364, is given in AAB47054. PRO364 PRO175 may be used in a mixture with a cardiovascular, endothelial,
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                                                                                                                                antiarthritic; antiinflammatory; antianaemic; immunosuppressive;
antithyroid; antidiabetic; neuroprotective; hepatotropic; viruci
dermatological; antipsoriatic; antiasthmatic; antiallergic;
                                                                                                                                                                                                                                                                                                         Human immunostimulant
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                                                                                                                                                                                                                                       PRO175; UNQ149; human;
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                                                                                                                                                                                                                                                                                                         PRO175
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Pred. No. 4.7e-97;
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Query Match Best Local Similarity

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951; DB 4; No. 4.7e-97;

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Sequence 177

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Cerythema multiforme and psoriasis), allergic disease, (auto) immune-mediated skin diseases (such as pullergic rythema multiforme and psoriated disease, (corythema multiforme and psoriated disease, (corythema multiforme and psoriatis), corythema multiforme and multiple sclerosis, idiopathic, corythema multiple sclerosis), autoimmune chronic corythyroiditis, diabetes mellitus, immune-mediated renal disease, corythyroiditis, diabetes mellitus, immune-mediated renal disease, corythyroiditis, primary biliary cirrhosis, granulomatous hepatitis, corythyrosis, granulomatous hepatitis, colitics and crohn's diseases, (such as bullous skin disease, corythyrosis, allergic diseases (such as sullergic diseases, such as sullous skin disease, corythyrosis, allergic diseases (such as asthma, corythyrosis, allergic diseases (such as asthma, corythyrosis, allergic diseases (such as sullous skin diseases, corythyrosis, allergic diseases (such as asthma, corythyrosis, allergic diseases (such as asthma, corythyrosis, allergic diseases (such as sullous skin diseases, corythyrosis, allergic diseases (such as asthma, corythyrosis, allergic diseases (such as sullous, asthma, corythyrosis, allergic diseases, such as sullous, and the corythyrosis, and corythyrosis, allergic diseases (such as sullous, and the corythyrosis, and a method of stanulous corporation and graft-versus-host collimed composition of the propertion, and graft-versus-host collimed composition of the propertion, and graft-versus-host collimed composition of the propertion
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Gurney AL, n.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             into a tissue of a mammal, stimulating or enhancing an immune response in a mammal, or increasing the proliferation of T-lymphocytes in a mammal in response to an antigen. Claimed compositions comprising the PRO polypeptide or its antagonist have the opposite effect. A claimed method for treating an immune related disorder, such as a T cell disorder, involves administering the PRO polypeptide, an agonist antibody or an antagonist antibody. The disorder is selected from systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (see AAF30050-62) encoding novel human PRO proteins (see AAB20108-20) including PRO175. Claimed compositions comprising these proteins or the agonists are useful for increasing infiltration of inflammatory cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New PRO polypeptides, nucleic acids and (ant)agonists, diagnosing and treating immune-related disorders, such sclerosis, rheumatoid arthritis and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is that of PRO175 (UNQ149), a novel human immunomodulator protein (20 kDa, pI 8.08) showing homology to tumour necrosis factor family members. The invention provides polympleteotides
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Hillan KJ, M
K, Wood WI;
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129. .133
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RESULT 5
AAB53065
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                                                                                                                                                                                                                                       01-SEP-1999;
08-SEP-1999;
15-SEP-1999;
15-SEP-1999;
05-OCT-1999;
                                            New isolated nucleic acid for producing a PRO polypeptide, analyzing genetic disorders and treating cardiovascular, endothelial or angiogenic
                                                                                                                    Ashkenazi AJ, Baker KP, Ferra
Godowski PJ, Gurney AL, Hilla
Paoni NF, Pitti RM, Watanabe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; angiogenesis-associated protein; PRO; endothelial cell growth; cardiac hypertrophy; cardiovascular disorder; endothelial disorder; angiogenic disorder; atherosclerosis; osteoporosis; hypertension; myocardial infarction; diabetic retinopathy; rheumatoid arthritis; Crohn's disease; psoriasis; endometriosis; ulcer; wound healing; cancer; Alzheimer's disease; Huntington's disease; stroke; drug screening;
                                                                                                                                                                                                                                                                                                     26-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                   30-NOV-1999
                                                                                                                                                                                                                                30-NOV-1999
                                                                                                                                                                                                                                                                                                                  20-JUL-1999
                                                                                                                                                                                                                                                                                                                                         02-JUN-1999
                                                                                                                                                                                                                                                                                                                                                               12-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                    05-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human angiogenesis-associated protein PRO175, SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB53065
                                                                                                                                                                   (GETH )
                                                                                  2001-090793/10.
DB; AAC97374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapy;
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                                                                                                           PU,
Pitti
                                                                                                                                                                   GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; protein; 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KEGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN 120
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                                   such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                    2000WO-US000219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                99WO-US020111.
99WO-US020594.
99WO-US021090.
99WO-US021547.
99WO-US023089.
99WO-US028313.
99WO-US028409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transgenic animal
                                                                                                                                                                                                                                                                                                     99US-
                                                                                                                                                                                           99WO-US028564.
99WO-US028565.
                                                                                                                                                                                                                                                                                                                 99US-0144758P.
                                                                                                                                                                                                                                                                                                                                       99WO-US012252.
                                                                                                                                                                                                                                                                                                                                                               99WO-US005028.
99US-0123957P.
                                atherosclerosis,
                                                                                                                                                                                                                                                                                                     -0145698P.
                                                                                                                                                                                                                                                                                                                                                     -0134287P.
                                                                                                                                                                                                                                                                                                                            -0141037P.
                                                                                                                   Ferrara N, Gerber H,
Hillan KJ, Kuo SS, M;
anabe CK, Williams PM,
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                                wounds or
                                  cancer.
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                                                                                                                               Mark MR,
                                                                                                                                         Goddard A;
                                                                                                                    Wood WI;
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                                                                                                                               Marsters SA;
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Claim 69; Fig

4; 293pp; English

Peptide

Location/Qualifiers

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RESULT 6
AAB47287
IID AAB47287
XX AAB4
XX AAB4
XX PRO1
DE PRO1
XX PRO1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                           PRO; type II transmembrane protein; tumour necrosis factor; stroke; heart hypertrophy; cardiovascular; endothelial; angiogenic; disorder; myocardial infarction; cardiac hypertrophy; PGF 2alpha; trauma; bone; cancer; age-related macular degeneration; wound; burn; hypertension; diabetes mellitus; osteoporosis; ischaemia, atherosclerosis; psoriasis; rheumatoid arthritis; Crohn's disease; amylotrophic lateral sclerosis; endometriosis; angina; neoplasms; periodontal disease; cartilage; TNF; Alzheimer's disease; Parkinson's disease; Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB47287 standard; protein; 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression or activity; diagnosing a cardiovascular, endothelial or angiogenic disorder, or a susceptibility to such a disorder by detecting mutations in a PRO gene, or the expression level of a PRO gene within a particular tissue; treating a cardiovascular, endothelial or angiogenic disorder via the administration of a PRO protein, PRO nucleic acid, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein, PRO antibodies specific for a PRO protein, fusion proteins comprising a PRO protein, agonists or antagonists of a PRO protein compounds which inhibit the expression of a PRO gene. The invention additionally encompasses methods of identifying modulators of PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to novel human angiogenesis-associated proteins designated PRO proteins (AAB53064-B53097), and to nucleic acids encode PRO proteins. The invention also relates to vectors and host cells comprising a PRO nucleic acid, the recombinant production of a PRO
                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRO175 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-AUG-2001
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Pred. No. 4.7e-97;
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                                                                                                                                                                                                The sequences given in AAB47287-90 show PRO polypeptides. PRO175 shows CC type II transmembrane protein typology, and portions of PRO364 show CC homology to members of the tumour necrosis factor (TNP) family, thereby indicating that it may be a novel member of the TNP family. PRO175 and CC PRO185 stimulated heart hypertrophy. PRO CDNA's may be used to identify a compound that inhibits PRO, diagnosing a cardio- vascular, endothelial or angiogenic disorder in a mammal by detecting PRO cDNA, treating CC cardiovascular, endothelial or angiogenic disorder in a mammal, and CC cardiovascular, endothelial or angiogenic disorder in a mammal, and CC cardiovascular in a mammal. The mammal is a human which has suffered myocardial infarction, cardiac hypertrophy or inhibiting endothelial cell growth or CC anglogenesis in a mammal. The mammal is a human which has suffered myocardial infarction, cardiac hypertrophy characterized by the presence of elevated PGF 2alpha, trauma, cancer or age-related macular degeneration. Trauma includes wounds or burns. Other treatable diseases include diabetes mellitus, osteoporosis, ischaemia, hyper- tension, crhoumatoid arthritis, Crohn's disease, atherosclerosis, psoriasis, conditional cardioral and cartilage repair, Alzheimer's disease, parkinson's disease, Huntington's disease, amaletrophic amaletrophic cardioral disease, Huntington's disease, Huntington's disease, amaletrophic cardioral disease, amaletrop
                                                                          Matches
                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JAN-2000; 2000WO-US000219.
24-FEB-2000; 2000WO-US005004.
15-MAR-2000; 2000WO-US006884.
30-MAR-2000; 2000WO-US008439.
17-MAY-2000; 2000WO-US013705.
30-MAY-2000; 2000WO-US014941.
28-JUL-2000; 2000WO-US020710.
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Hillan KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cardiovascular, endothelial or an mellitus, myocardial infarction,
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30-NOV-1999;
                                                                                                                                                  Sequence
                                                                                                                                                                                      amylotrophic lateral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56; Fig 2; 144pp; English.
                                                                      al Similarity
177; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENENTECH
                                                                                                                                                    177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRO polypeptide useful in treating and dia scular, endothelial or angiogenic disorder, myocardial infarction, arthritis.
                    MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
	exttt{MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA}
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                                                                      100.0%; llarity 100.0%; Conservative 0
                                                                                                                                                  AA;
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99WO-US028409.
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/label
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/label= Transmembrane
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/label= N-myristoylation
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                                                                                                                                                                                      sclerosis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-linked glycosylation site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Extracellular domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Signal peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ferrara N, Godowsk
sters SA, Paoni NF,
                                                                      0;
                                                                    Score 951; DB 4;
Pred. No. 4.7e-97;
; Mismatches 0;
                                                                                                                                                                                        and stroke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              glycosylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Godowski PJ, (
aoni NF, Pitti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l diagnosing
der e.g. can
                                                                                                        Length 177;
                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JAN-2000;
11-FEB-2000;
18-FEB-2000;
18-FEB-2000;
24-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUL-1999;
26-JUL-1999;
28-JUL-1999;
01-SEP-1999;
30-NOV-1999;
30-NOV-1999;
02-DEC-1999;
16-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; PRO; cardiant; antiangiogenic; antiarteriosclerotic; hyasotropic; antirheumatic; antiarthritic; antiinflammatory; c vulnerary; antianginal; gene therapy; cardiovascular disease; endothelial disorder; angiogenic disorder; cancer; periodonta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB50979 standard;
                                                                                                                                                                                                                                                                                                                           Ashkenazi AJ, Baker
Goddard A, Godowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-MAR-2000;
10-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUN-1999;
23-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO175 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB50979;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200073445-A2
                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           healing.
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                                                                                                                                                                                                                                                                                                                  A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                        2000WO-US0030095

2000WO-US000319

2000WO-US0003565

2000WO-US003565

2000WO-US004341

2000WO-US004342

2000WO-US005004

2000WO-US005004

2000WO-US00500819

2000WO-US006819

2000WO-US007532

2000WO-US008439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US020111.
99WO-US028313.
99WO-US028409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0145698P
99US-0146222P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US028565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US012252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein;
                                                                                                                                                                                                                                                                                                                           일 등
                                                                                                                                                                                                                                                                                                 KP, Ferrara N,
PJ, Gurney AL,
Watanabe CK, W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              177
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                                                                                                                                                                                                                                                                                                   , Gerber H, Gerritsen
, Kuo SS, Mark MR, Ma
Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   periodontal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypotensive;
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The present sequence is one of seventeen novel PRO polypeptides. nucleic acids, polypeptides, agonists and antagonists are useful treating cardiovascular, endothelial or angiogenic disorders in a

in a

mammal

The PRO

Seventeen nucleic acids diagnosis and treatment

nucleic acids encoding PRO polypeptides which are useful in and treatment of cardiovascular, endothelial or angiogenic in a mammal.

disorders

Claim

71; Fig 2; 182pp; English.

WPI; 2001-025251/03. N-PSDB; AAC90563.

Paoni

Pitti RM,

Marsters

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
 18-FEB-2000;
22-FEB-2000;
24-FEB-2000;
02-MAR-2000;
15-MAR-2000;
17-MAY-2000;
                                                                                                    30-NOV-1999;
01-DEC-1999;
05-JAN-2000;
06-JAN-2000;
11-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antagonists are also used to prevent tumour anglogenesis and for treating periodontal diseases. They are also used to stimulate wound healing and tissue regeneration. The PRO nucleic acids, polypeptides and anti-PRO antibodies are useful for diagnosing a cardiovascular, endothelial or anglogenic disorder
                                                                                                                                                                              15-SEP-1999;
29-NOV-1999;
                                                                                                                                                                                                            28-JUL-1999;
13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             Human; PRO; antiinflammatory; ophthalmological; vasotropic; retinal cell injury; ocular disease; retinitis pigmentosa; macular degeneration; retinal detachment; retinal tear; retinopathy; retinal degenerative disease; macular hole; degenerative myopia; acute retinal necrosis syndrome; traumatic chorioretinopathy; purtscher's retinopathy; oedema; ischaemic condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Examples of these disorders include cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restenosis, Reynaud's disease, rheumatoid arthritis, angina, mycardial infarctions, thrombophlebitis and lymphangitis. The PRO polypeptides and
                                                                                                                                                                                                                                                                                     08-FEB-2001.
                                                                                                                                                                                                                                                                                                                  WO200109327-A2.
                                                                                                                                                                                                                                                                                                                                                                                        Purtscher's retinopathy; oedema; ischaemic condition; retinal vision occlusion; collagen vascular disease; thrombocytopaenic purpura; uveitis; retinal vasculitis; Eales
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU81953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU81953 standard; protein; 177
                                                                                                                                                                                                                                                      28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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177; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MCLSHLENMPLSHSRTQGAQRSSWKLMLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
                                                                                                                                                                                                                                                                                                                                                                          lupus erythematosus; environmental trauma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              177
2000WO-US000219
2000WO-US000376
2000WO-US003565
2000WO-US004411
2000WO-US00504
2000WO-US00504
2000WO-US0050841
2000WO-US005841
2000WO-US006884
2000WO-US00843
2000WO-US00843
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                                                                                                                                                                                                                                                       2000WO-US020710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                 99WO-US028313.
99WO-US028301.
                                                                                                                                                                              99WO-US028214.
                                                                                                                                                                                                99WO-US021090.
                                                                                                                                                                                                            99WS-0146222P.
99WO-US020944.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 951; DB 4;
Pred. No. 4.7e-97;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for treating
                                                                                                                                                                                                                                                                                                                                                                                         disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                               encoding the PRO proteins, a vector comprising the nucleic acid, a host coell comprising the vector, and anti-PRO antibody. The PRO proteins are useful for promoting survival of retinal cells (retinal neurons such as retinal ganglion cells, displaced retinal ganglion cells, amacrine cells, or supportive cells such as Muller cells or pigment cepthelial cells), or delaying or preventing retinal cell injury or death caused by ocular disease (which is or is associated with retinitis pigmentoss, macular degeneration, retinal detachment, retinal tear, cretinopathy, retinal degenerative disease, macular hole, degenerative myopia, acute retinal necrosis syndrome, traumatic chorioretinopathy or contusion, purtscher's retinopathy, oedema, an ischaemic condition, central or branch retinal vision occlusion, collagen vascular disease, contusion branch retinal vision occlusion, central or branch retinal vision occlusion vasculatitis, occlusion associated with Eales disease or systemic lupus erythematosus), retinal call injury or death is condition, the condition of the condition occlusion.
                                                                                                                                                                           Matches
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                             Sequence 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to promoting the survival of retinal cells, or delaying or preventing retinal cell injury or death, by contacting the retinal cells with the polypeptide such as PRO175, PRO200, PRO216, PRO241, PRO306, PRO346, PRO322, PRO536, PRO943, PRO840, PRO828, PRO826, PRO1068 or PRO1132 polypeptide. Also included are the nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 44; Fig 2; 152pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Promoting survival of retinal cells, or delaying or preventing retinal cell injury or death, by contacting retinal cells with PRO175, 220, 216, 243, 306, 346, 322, 536, 943, 840, 828, 826, 1068 or PRO1132 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB;
                                                                                                                                                                                                                                                                                                         delayed or prevented by substantially not causing angiogenesis or mitogenesis. The present sequence represents a PRO protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kljavin IJ,
Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ashkenazi AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH
                                                                                                                                                                                               Local
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61
                                           61
                                                                                                                               Н
                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK28562.
                             KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN 120
KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN 120
                                                                                                                               MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA 60
                                                                                      MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                               AĄ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , Baker KF,
Lafleur M,
                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Goddard A, Godows
Mark MR, Marsters
                                                                                                                                                                           0;
                                                                                                                                                                                             Score 951; DB 5;
Pred. No. 4.7e-97;
                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Godowski PJ, Gur
ırsters SA, Pitti
                                                                                                                                                                           0
                                                                                                                                                                                                                  Length 177;
                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gurney AL;
tti RM, Wa
                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Watanabe CK;
                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or death
                                                                                      60
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RESULT 9 ABR42323

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KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS

KDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS

121 121

ABR42323 standard; protein; 177 AA

SXXXXXXXXXXXXXX ABR42323;

(first

Human GITRL protein.

Human; GITRL; tumour necrosis factor; ligand;

immunomodulator; osteopathic.

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RESULT 10
ABP60544
ID ABP6054A
AC ABP60
XX ABP60
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KW APRIL
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Best Local S
Matches 177
              dermatological; immunosuppressive; antiinflammatory; antirheumatic; antiarthritic; cytostatic; antianaemic; antiallergic; antiasthmatic; neuroprotective; ophthalmological; tuberculostatic; antidiabetic; antipsoriatic; anti-HIV; antiarteriosclerotic; vasotropic; thyromimetic; haemostatic; cancer; autoimmune disease; graft versus host disease; GVHD; inflammatory disorder; proliferative disorder; single chain antibody; antibody; human; tumour necrosis factor; endokine-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to compositions comprising heterotrimeric of tumour necrosis factor (TNF) ligand family members, and their the detection, prevention and treatment of disease. In preferred embodiments, the heterotrimeric complex comprises full-length or extracellular portions of GITRL together with full-length or extracellular portions of other TNF ligand family members. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New heteromultimeric complex having a first polypeptide member of tumor necrosis factor (TNF) ligand family, and a second different of TNF ligand family, useful for treating cancer, osteoporosis or
                                                                                                                                                                            Human tumour necrosis factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                heterotrimeric complexes of the invention are useful for treating autoimmune disease, cancer or osteoporosis, and particularly for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                  ABP60544 standard;
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DB; ACC57909.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA;
                                                                                                                                       immunospecific; tumour necrosis factor delta; TNF-delta;
                                                                                                                                                                                                           (first entry)
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No. 4.7e-97;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel antibody or its fragment, which immunospecifically binds tumour necrosis factor Delta (TNF-delta/APRIL). The antibody of the invention has dermatological, immunosuppressive, antiinflammatory, antixheumic, antiarthritic, cytostatic, antianaemic, antiallergic, antiasthmatic, neuroprotective, ophthalmological, tuberculostatic, antiidiabetic, antipsoriatic, anti-HIV, antiarteriosclerotic, vasotropic, thyromimetic, and haemostatic activity. The antibody or its fragment are useful for treating, preventing or ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in human, disease or disorder such as autoimmune disease, and graft versus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APRIL receptor expression or aberrant function of APRIL or APRIL receptor. The disease or disorders includes autoimmune and inflammatory receptor. The disease or disorders includes autoimmune neutropenia, haemolytic anaemia, dermatitis, asthma, allergic encephalomyelitis, myocarditis, multiple sclerosis, uveitis, tuberculosis, diabetes mellitus, psoriasis, cancer of the immune system, particularly B cell cancers, immune disorders such as myasthenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          host disease (GVHD). The autoimmune disease is systemic lupus erythematosus, rheumatoid arthritis or Sjogren's syndrome. The antibody is useful for detecting, diagnosing, prognosing, treating, preventing or suseful for detecting, diagnosing, prognosing, treating, preventing or manufactured with aberrant APRIL or ameliorating a disease or disorder associated with aberrant APRIL or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        infectious diseases (e.g. acquired immunodeficiency syndrome (AIDS)), proliferative disorders (e.g. leukemia). The present sequence represe the tumour necrosis factor endokine-alpha
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Hodgkin's lymphoma, m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 214-215;
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multiple myeloma,
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ma, rheumatoid arthritis or Sjogren's
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No. 4.7e-97;
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ADC35222; ADC35222

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(first entry)

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Best Local
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                                  ABW02285;
                                                                                            ABW02285 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New Endokine alpha gene useful for preparing a composition for treating disease associated with excessive or insufficient bone resorption e.g., osteoporosis, Paget's disease or arterial calcification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 40; 145pp; English.
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30-OCT-2001;
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100.0%; Pred. No. 4.7e-97;
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                                                                                                                                                                                                      ameliorating a disease or disorder of the gastrointestinal tract. The method involves administering a composition comprising tumour necrosis factor (TNF)-gamma-beta and its receptors DR3 and TR6 to a person with, or suspected of having the disease or disorder. The antagonist of TNF-gamma-beta is useful for treating or ameliorating a gastrointestinal tract disease or disorder, specifically an inflammatory bowel disease, e.g. Crohn's disease or ulcerative colitis. The present sequence is human endokine alpha protein. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-2001;
06-JUL-2001;
24-AUG-2001;
07-DEC-2001;
23-AUG-2002;
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30-APR-1999;
03-MAY-1999;
13-MAY-1999;
18-FEB-2000;
27-APR-2000;
07-JUL-2000;
                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 153-154; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                              Use of tumor necrosis factor gamma-beta antagonists for treating of ameliorating a disease or disorders of the gastrointestinal tract, inflammatory bowel disease, Crohn's disease or ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB;
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07-AUG-1998;
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09-JAN-1998;
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DB; AAD63921.
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                                                                            MCLSHLENMPLSHSRTQGAQRSSWKLMLFCSIVMLLFLCSESWLIFIFLQLETAKEPCMA 60
KDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS
                                             KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN
                                                                                                   MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA 60
                         KFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKN
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                                                                    This polypeptide comprises human endokine-alpha, a novel member of the tumour necrosis factor (TNF) family of cytokines that shows 30% similarity and 22% identity to human TNF-alpha. Its amino acid sequence was deduced from a human striatum cDNA clone (see AAV19195). Isolation of this nucleic acid allows production of recombinant endokine-alpha polypeptides in transformed host cells. Endokine-alpha may be involved in disorders of immunomodulation, infection, cell proliferation,
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angiogenesis, tumour metastasis, apoptosis, sepsis and endotoxaemia, and may be useful for treating melanoma and sarcoma, particularly where coupled to a radioisotope or cytostatic agent. Antibodies raised against
                                                                                                                                                                                                                                                                                               Nucleic acid
treatment of
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                                                                                                                                                                                                                                                                                             encoding human endokine-alpha - useful for diagnosis tumour necrosis factor-related diseases.
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KW autoimmune disease; lupus; multiple sclerosis; chronic inflammation;
KW acute inflammation; acute allograft rejection; graft versus host disease;
KW transplant rejection; foetal resorption; faecal peritonitis; allergy;
KW bowel disease; sepsis; leukaemia; chronic hypergammaglobulinemeia;
KW bowel disease; sepsis; leukaemia; chronic hypergammaglobulinemeia;
KW downondritis; scleroderma; Wegener granulomatosis; dermatomyositis;
KW chronic active hepatitis; myasthenia gravis; psoriasis; vitilogo;
KW cteven-Johnson syndrome; idiopathic sprue; gluten-sensitive enteropathy;
KW gemphigus vulgaris; Goodpasture's disease; bullous pemphigoid;
KW discoid lupus; dense deposit disease; endocrine ophthalmopathy;
KW irritable bowel disease; asthma; Grave's disease; sarcoidosis; cirrhosis;
KW juvenile diabetes; insulin dependent diabetes mellitus; uveitis;
KW lymphopenias; polyarteritis nodosa; Sjogren's syndrome; Bechet's disease;
KW primary myxedema; polymyositis; mixed connective tissue disease;
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                                                                     16-AUG-1996;
                                                                                                                                                                                                                07-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                      Human; endokine alpha; tumour necrosis factor; TNF; cytokine; immunomodulation; inflammation; cell proliferation; angiogene tumour metastasis; apoptosis; sepsis; endotoxemia.
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   (HUMA-) HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT 128
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Pred. No. 4.4e-92;
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                                                                                                                                                                                                                                                 Sequence 169
                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Fig 1; 31pp; English.
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                                                                                                                                             MPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSK 68
                                                              WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT
                                                                                WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT 128
                                                                                                                         MPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSK 60
NKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS
                                                                                                                                                                                      Conservative
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                                                                                                                                                                                    95.3%; Score 906; DB 3; I
100.0%; Pred. No. 4.4e-92;
Ltive 0; Mismatches 0;
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169
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Search completed: November Job time: 159 secs 2, 2004, 12:37:39

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Database
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seq length: 0
seq length: 2000000000
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                                                                                                                                                                                                                                  1 MCLSHLENMPLSHSRTQGAQ......VLKNNTYWGIILLANPQFIS 177
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Copyright (c) 1993 - 2004 Compugen Ltd.
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/cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

13	12	11	10	9	8	7	o,	ហ	4	ω	N	н	Result No.
79.5	85.5	281	608	809	906	906	951	951	951	951	951	951	Score
8.4	9.0	29.5	63.9	63.9	95.3	95.3	100.0	100.0	100.0	100.0	100.0	100.0	Query Match Length
246	377	52	113	113	169	169	177	177	177	177	177	177	Length
15	14	9	16	14	13	9	15	14	14	14	13	13	BB
US-10-424-599-201262	US-10-295-027-280	US-09-864-761-48455	US-10-611-399-17	US-10-338-083-17	US-10-136-511-2	US-09-345-790-2	US-10-202-062-40	US-10-310-793-44	US-10-218-547-40	US-10-151-882-39	US-10-116-378-16	US-10-080-455-1	ID
Sequence 201262,	Sequence 280, App	Sequence 48455, A	Sequence 17, Appl	Sequence 17, Appl	Sequence 2, Appli	Sequence 2, Appli	Sequence 40, Appl	44,	Sequence 40, Appl	Sequence 39, Appl	Sequence 16, Appl	Sequence 1, Appli	Description

ALIGNMENTS

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publication No. US20020146389A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Godowski, Paul J.
                                                                                             Query Match 100.0%; Score 951; DB 13; Best Local Similarity 100.0%; Pred. No. 2e-92; Matches 177; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                         APPLICANT: Gurney, Austin I.
APPLICANT: Mark, Melanie R.
APPLICANT: Mark, Melanie R.
APPLICANT: Mark, Melanie R.
APPLICANT: Markers, Scot A.
APPLICANT: Pitti, Robert M.
TITLE OF INVENTION: DNA19355 Polypeptide, A Tumor Necrosis Factor Homolog
FILE REFERENCE: P1150R2
CURRENT APPLICATION NUMBER: US/10/080,455
CURRENT FILING DATE: 1998-11-18
PRIOR APPLICATION NUMBER: US/0/69,661
PRIOR APPLICATION NUMBER: US 60/065,635
PRIOR FILING DATE: 1997-11-18
PRIOR APPLICATION NUMBER: US 60/065,635
PRIOR FILING DATE: 1997-11-18
                                                                                                                                                                                                             LENGTH: 177
TYPE: PRT
ORGANISM: Homo sapiens
1 MCLSHLENMPLSHSRTQGAQRSSWKLMLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Baker, Kevin P.
Godowski, Paul J.
Gurney, Austin L.
Mark, Melanie R.
                                                                                                                                          Length 177;
                                                                                                Indels
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                                                                                             Gaps
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US-10-151-882-39

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RESULT 3
US-10-151-882-39
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APPLICANT: Ashkenazi, A
APPLICANT: Goddard, Au
APPLICANT: Gurney, Aus
APPLICANT: Marsters, S

                                                                                                                          CURRENT FILING DATE: 2002-05-22
PRIOR APPLICATION NUMBER: 60/293,100
PRIOR FILING DATE: 2001-05-24
NUMBER OF SEQ ID NOS: 48
SOFTWARE: PatentIn version 3.0
SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 39, Application US/10151882
Publication No. US20030059862A1
GENERAL INFORMATION:
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Antibodies Against Tumor Necrosis Factor Delta (APRIL)
FILE REFERENCE: PF554
CURRENT APPLICATION NUMBER: US/10/151,882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: P1206R1
CURRENT APPLICATION NUMBER: US/10/116,378
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/247,225
PRIOR FILING DATE: EARLIER FILING DATE: 1999-02-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/074,0
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-09
NUMBER OF SEQ ID NOS: 31
SEQ ID NO 16
LENGTH: 177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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APPLICANT: Gurney, Austin
APPLICANT: Marsterr, Scot A.
APPLICANT: Mod, William
APPLICANT: Wood, William
TITLE OF INVENTION: NOVEL TUMOR NECROSIS FACTOR RECEPTOR HOMOLOG AND
TITLE OF INVENTION: ACIDS ENCODING THE SAME
ORGANISM: Homo sapiens
                                                                                                 ENGTH: 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             177;
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RESULT 5
US-10-310-793-44
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                                                                                                                                                                                         Sequence 44, Application US/10310793 Publication No. US20030198640A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local Similarity
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Best Local Similarity
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               APPLICANT: Rosen, Craig A
APPLICANT: Zhang, Jun
APPLICANT: Wei, Ping
TITLE OF INVENTION: Methods And Compositions |
TITLE OF INVENTION: Relating To Human Tumor |
FILE REFERENCE: PF573
                                                                                                                                                 APPLICANT: Yu, Guo-Liang APPLICANT: Ni, Jian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/218,547
CURRENT FILING DATE: 2002-08-15
PRIOR APPLICATION NUMBER: 60/312,542
PRIOR FILING DATE: 2001-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Methods And Compositions TITLE OF INVENTION: Human Endokine Alpha FILE REFERENCE: PF561
CURRENT APPLICATION NUMBER: US/10/310,793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/2 PRIOR FILING DATE: 2001-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 177
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                   121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  177;
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Pred. No. 2e-92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 14;
                                         Necrosis Factor-Gamma
                                                             For Treating Inflammatory Bowel Diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     For Treating Metabolic Bone Diseases
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PRIOR APPLICATION NUMBER: 60/336,695 PRIOR FILING DATE: 2001-12-07

CURRENT FILING DATE:

2002-12-06

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US-10-202-062-40
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                                                                                                                                      US-10-202-062-40
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SEQ ID NO 44
LENGTH: 17
                                                                                                                                                                                                                 SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Publication No. US20040038349A1
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                                                        Best Local Similarity Matches 177; Conserv
                                                                               Query Match
Best Local :
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NUMBER OF SEQ ID NOS: 71
COFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/202,062
CURRENT FILING DATE: 2002-07-25
PRIOR APPLICATION NUMBER: 60/307,838
PRIOR FILING DATE: 2001-07-27
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PRIOR FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 60/216,879
PRIOR APPLICATION NUMBER: 60/216,879
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PRIOR FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: 60/314,381
PRIOR FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: 09/899,059
PRIOR FILING DATE: 2001-07-06
PRIOR FILING DATE: 2001-07-06
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Human Genome Sciences, Inc.,
TITLE OF INVENTION: Heteromultimeric TNF Ligand Family members
                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn versi
                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PF559
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                                                                                                                                                        TYPE: PRT ORGANISM: human
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                                                                                                                                                                                             LENGTH: 177
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FILING DATE: 1999-05-13
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APPLICATION NUMBER: 60/180,908
FILING DATE: 2000-02-08
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MCLSHLENMPLSHSRTQGAQRSSWXLMLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA 60
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                                                                                                                                                                                                                                     version 3.0
                                                                             100.0%;
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100.0%; Pred. No. 2e-92;
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                                                      Score 951; DB 15;
Pred. No. 2e-92;
Mismatches 0;
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Patent No. US20020099198A1
GENERAL INFORMATION:
APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig A.
                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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ZIP: 2005-395-
ZIP: 2005-395-
COMPUTER READABLE FORM:
COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
CO
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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NAME: Steffe, Exic K.
REGISTRATION UNMER: 36,688
REFERENCE/DOCKST NUMBER: 1488
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
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TITLE OF INVENTION: Human Endokine Alpha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
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                                        129 NKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 177
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                                                                                                                                       WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT
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                                                                                                                                                                                                                                                                                                                    MPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSK 68
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                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    95.3%; Score 906; DB 9;
100.0%; Pred. No. 1.1e-87;
tive 0; Mismatches 0;
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US-10-136-511-2

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US-10-338-083-17
; Sequence 17, Application US/10338083
; Publication No. US20030166559A1
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                                                     ; ORGANISM: Homo sapiens
US-10-338-083-17
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PRIOR FILING DATE: 2000-02-25
PRIOR PELICATION NUMBER: 09/345,790
PRIOR FILING DATE: 1999-07-01
PRIOR FILING DATE: 1999-05-28
PRIOR APPLICATION NUMBER: 60/122,099
PRIOR FILING DATE: 1999-02-26
PRIOR FILING DATE: 1999-02-26
PRIOR APPLICATION NUMBER: 60/122,099
PRIOR APPLICATION NUMBER: 08/912,227
PRIOR APPLICATION NUMBER: 60/024,058
PRIOR FILING DATE: 1996-08-16
NUMBER OF SEQ ID NOS: 10
                                                                                                     SOFTWARE: PatentIn version 3.2
SEQ ID NO 17
LENGTH: 113
TYPE: PRT
  Query Match
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LENGTH: 169
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Publication No. US20020168729A1
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Best Local &
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                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/338,083
CURRENT FILLING DATE: 2003-01-16
PRIOR APPLICATION NUMBER: US 60/345,805
PRIOR FILLING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR FILLING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Tansey, Malu Lourdes G.
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: Dominant Negative Proteins and Methods Thereof
FILE REFERENCE: A-71273-2
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CURRENT APPLICATION NUMBER: US/10/136,511
CURRENT FILING DATE: 2002-05-02
PRIOR ADDITORNOON
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APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Human E
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les 169; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NKSKIQNVCGTYELHVCDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95.3%; Score 906; DB 13; Length 169; ilarity 100.0%; Pred. No. 1.1e-87; Conservative 0; Mismatches 0; Indels
  63.9%;
  Score 608;
DB 14;
  Length 113;
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; TYPE: PRT; ORGANISM: Homo sapiens US-10-611-399-17
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US-10-611-399-17
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CURRENT FILING DATE: 2003-07-01
PRIOR APPLICATION NUMBER: US 10/338,083
PRIOR FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/345,805
PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 23
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APPLICANT: Desjariais, John R.
APPLICANT: Tansey, Malu Lourdes G.
APPLICANT: Dahiyat, Bassil I.
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: DOMINANT NEGATIVE PROTEINS AND METHODS THEREOF FILE REFERENCE: A-71273-3
                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL
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APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeomica-X-1
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PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                     APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K
APPLICANT: Chen, Wensheng
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; Mismatches 0;
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FOR

APPLICATION NUMBER: US 60/236,359

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RESULT 12
US-10-295-027-280
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Best Local S
Matches 52
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                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                       APPLICANT: Murray, Richard
APPLICANT: Watson, Susan R.
APPLICANT: Eos Biotechnology, Inc.
TITLE OF INVENTION: Methods of Diagnosis of Cancer, Compositions and
TITLE OF INVENTION: Methods of Screening for Modulators of Cancer
FILE REFERENCE: 018501-012500US
CURRENT APPLICATION NUMBER: US/10/295,027
CURRENT FILING DATE: 2002-11-13
CURRENT FILING DATE: 2002-11-13
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FERATURE:
OTHER INFORMATION: MAP TO AL031599.1
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.2
OTHER INFORMATION: EST_HUMAN HIT: BE270854.1, EVALUE 6.50e-01
OTHER INFORMATION: SWISSPROT HIT: P25003, EVALUE 1.20e+00
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APPLICATION NUMBER: US 09/663,733 FILING DATE: 2000-09-15 APPLICATION NUMBER: US 60/350,666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 09/774,203 FILING DATE: 2001-01-29
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APPLICATION NUMBER: PCT/US01/00664
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APPLICATION NUMBER: PCT/US01/00666
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Mack, David H.
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100.0%; Pred. No. 3.
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US-10-424-599-201262
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US-10-424-599-201262
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                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 201262
LENGTH: 246
                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 201262, Application US Publication No. US20040031072A1
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Best Local
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PRIOR FILING DATE: 2001-11-15
PRIOR FILING DATE: 2001-11-21
PRIOR FILING DATE: 2001-11-21
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PRIOR APPLICATION NUMBER: US 60/347,349
PRIOR FILING DATE: 2002-01-10
PRIOR APPLICATION NUMBER: US 60/355,250
PRIOR FILING DATE: 2002-02-08
PRIOR APPLICATION NUMBER: US 60/356,714
PRIOR FILING DATE: 2002-02-13
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NUMBER OF SEQ ID NOS: 1386
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PRIOR FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/347,211
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 2001-11-15
                                                                                                                        NAME/KEY: unsure
LOCATION: (1)..(246
OTHER INFORMATION:
                                                                                                                                                                                           TYPE: PRT
ORGANISM: Glycine max
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                                                                                       INFORMATION:
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ch 8.4%; l Similarity 22.9%; 36; Conservative 2
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Cao Yongwei
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Score 79.5; DB
Pred. No. 6.1;
21; Mismatches
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Pred. No. 2.5;
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US-10-149-310-52
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Best Local Similarity
Matches 37; Conserv
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 52
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Publication No. US20040077039A1
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APPLICANT: Madden, Kevin T.
APPLICANT: Madden, Kevin T.
APPLICANT: Maxon, Mary
APPLICANT: Maxon, Mair
IITLE OF INVENTION: Modulation of Secondary Metabolite Production by
IITLE OF INVENTION: Modulation of Secondary Metabolite Production by
FILE REFERENCE: 14184-019US1
CURRENT APPLICATION NUMBER: US/10/149,310
CURRENT FILING DATE: 2003-02-19
PRIOR APPLICATION NUMBER: PCT/US01/29288
PRIOR FILING DATE: 2001-09-19
APPLICANT: Sherman, Amir
TITLE OF INVENTION: Modulation of Secondary Metabolite Production
TITLE OF INVENTION: Zinc Binuclear Cluster Proteins
FILE REFERENCE: 14184-019US1
CURRENT APPLICATION NUMBER: US/10/149,310
CURRENT FILLING DATE: 2003-02-19
PRIOR APPLICATION NUMBER: PCT/US01/29288
PRIOR FILING DATE: 2001-09-19
                                                                                                                                                         APPLICANT:
                                                                                                                                                                           APPLICANT: Holtzman, Douglas
APPLICANT: Madden, Kevin T.
APPLICANT: Maxon, Mary
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Pred. No. 32
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Search completed: November Job time: 130 secs
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SEQ ID NO 50
LENGTH: 880
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Matches
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PRIOR FILING DATE: 2000-09-19
NUMBER OF SEQ ID NOS: 308
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
                                                                                                                                                                                                                                                                                                   Local Similarity
les 37; Conserv
                                                                          672 -ENIIGTDSLYGLENSLILLFSDCVRIVRHNEYYNLTYLPVPR 713
                                                                                                                                                  615 ALNENDGKIHIEFVKEPITNVSADSTPSSTTPPIFTNIA---TESYYNKSDISKLVSKTD 671
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                                                                                                             133 IQNVGGTYELH-VGDTIDLIFNSEHQVLKNNTYWGIILLANPQ 174
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                     2, 2004, 12:52:46
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                     Score
    seq length: 0
seq length: 2000000000
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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    MCLSHLENMPLSHSRTQGAQ......VLKNNTYWGIILLANPQFIS
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    GenCore version 5.1.6 (c) 1993 - 2004 Compugen
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 US-08-912-227-2
US-09-345-790-2
US-09-107-532A-3959
US-09-107-532A-3959
US-09-248-796A-20671
US-09-248-796A-20671
US-08-089-397A-11
5395759-2
US-08-089-397A-11
5395759-2
US-09-0328-352-5760
US-09-248-352-5760
US-09-248-357-2
PCT-US-3-0474-5-6
US-09-248-796A-15058
US-09-248-796A-15058
US-09-248-796A-18800
US-09-248-796A-18800
US-09-248-796A-18800
US-09-248-933-10
US-08-515-903A-10
US-08-515-903A-10
US-08-515-903A-10
US-08-912-227-3
US-08-883-086-6
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Sequence 2, Appli
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Sequence 601, App
Sequence 20671, Appli
Sequence 11, Appli
Sequence 2, Appli
Sequence 3, Appli
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Sequence 6, Appli
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Matches

169; Conservative

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Mismatches

0;

Indels

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Gaps

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9 MPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSK

US-08-912-227-2 US-08-912-227-2 Sequence 2, Application US/08912227 Patent No. 5998171 GENERAL INFORMATION: Jon Liang APLICANT: No. Guo-Liang APLICANT: Ni., Guo-Liang APLICANT: Ni., Guo-Liang APLICANT: Ni., Jian APLICANT: Manufactory (CORRESPONDENCE ADDRESS: 10 CORRESPONDENCE ADDRESS: 10 CORRESPONDENCE ADDRESS: 10 CORRESPONDENCE ADDRESS: 10 COUNTRY: WASHINGTON STATE: DC COUNTRY: USA ZIP: 2006-3934 COUNTRY: USA ZIP: 2006-3934 COMPUTER: BEADABLE FORM: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: LBM PC Compatible COMPUTER: USA ZIP: 2006-3934 COMPUTER: USA ZIP: 2006-3934 COMPUTER: USA ZIP: 2006-3934 COMPUTER: HEADABLE FORM: MEDIUM TYPE: HEAPTON DATA: MEDIUM TYPE: USA ZIP: 2006-3934 COMPUTER: USA ZIP: 2006-3934 ZIP: 2006	ALIGNMENTS	28 71.5 7.5 233 3 US-08-880-342-37 Sequence 37, Appli 29 71.5 7.5 233 4 US-09-589-287B-3 Sequence 3, Appli 30 71.5 7.5 233 4 US-09-513-584-3 Sequence 3, Appli 31 71.5 7.5 233 4 US-09-137-864-9 Sequence 9, Appli 32 71.5 7.5 233 4 US-09-345-790-3 Sequence 5, Appli 33 71.5 7.5 233 4 US-09-345-790-3 Sequence 2, Appli 34 71.5 7.5 233 4 US-09-888-22 Sequence 3, Appli 36 71.5 7.5 233 4 US-09-888-947A-3 Sequence 3, Appli 38 71.5 7.5 233 4 US-09-588-947A-3 Sequence 3, Appli 39 71.5 7.5 233 4 US-09-589-286A-3 Sequence 3, Appli 39 71.5 7.5 233 4 US-09-589-286A-3 Sequence 3, Appli 41 71.5 7.5 233 4 US-09-005-874-3 Sequence 3, Appli 42 71.5 7.5 233 5 PCT US95-12840-10 Sequence 10, Appli 42 71.5 7.5 693 3 US-09-60-048-11 Sequence 11, Appl 44 71.5 7.5 693 3 US-09-880-866-11 Sequence 21, Appli 5 Sequence 21, App	
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                                                                       ; MOLECULE TYPE: protein US-09-513-584-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
Query Match 95.3%; Score 906; DB 4; Best Local Similarity 100.0%; Pred. No. 2.8e-99; Matches 169; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OCURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/513,584
FILING DATE: 25-FEB-2000
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/024,058
PRIOR APPLICATION NUMBER: US/09/024,058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Human Endokine Alpha and Methods
TITLE OF INVENTION: of Use
                                                                                                                                                                                                                                                                                                          FILING DATE: 01-JUL-1999
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/345,790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 15-AUG-
FILING DATE: 15-AUG-
PRIOR APPLICATION DATA:
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                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX. ADDRESSEE: P.L.L.C.
STREET: 1100 NEW YORK AVE., NW, SUITE 600
CITY: WASHINGTON
                                                                                                                                                                                   TELEPHONE: 202-0...
TELEPHONE: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 60/122,099 FILING DATE: 26-FEB-1999
                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 36,688 REFERENCE/DOCKET NUMBER: 14
                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/136,788 FILING DATE: 28-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 20005-3934
                                                                                                             TOPOLOGY:
                                                                                                                                             LENGTH:
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                                     DB 4; Length 169;
   Indels
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Gaps
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US-09-345-790-2
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                                                                                                                                                                                                                                          Best Local
                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT:
APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/345,790
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NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                             LENGTH: 169 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE
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Ni, Jian
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121
                           129 NKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANDQFIS 177
                                                                                                                                 69 WQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT
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NKSKIONVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 169
                                                                                                                                                                                           MPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSK
                                                                                             WOMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLT
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NW, SUITE 600
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RESULT 5
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US-09-107-532A-3959
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Patent No. 6583275
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
                Sequence 601, Application US/09538092
Patent NO. 6753314
GENERAL INFORMATION:
APPLICANT: Giot, Loic
APPLICANT: Mansfield, Traci A.
                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 3959:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...1
SEQUENCE DESCRIPTION: SEQ ID NO:
OF INVENTION: Protein-Protein Complexes and Method of Using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMÂTION:
NAME: ATINIELLO, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/107,532A FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                               515
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                                                                                                                                                                                                                                                                                                                      102
                                                                                                                                                                                                                                       157 QVLKNN 162
                                                                                                                                                                                                                                                                               APNGYQLDSSEFSITVKENQNVTTTRTNKKSTGSVEIEKIGDLDGLGLPNVVFTIYNSDN
                                                                                                                                                                                                                                                                                                                  APNANYNDVAPFEVRLYKNKDMIQTLTNKS-----KIQNVGGTYELHVGDTIDLIFNSEH
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                                                                                                                                                                                                  KVVKDN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: <Unknown> SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH: 1221 amino acids
                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                  580
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                                                                                                                                                                                                                                                                                                                                                                            8.9%; Score 84.5; DI
28.8%; Pred. No. 1.4;
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AND AMINO ACID SEQUENCES RELATING TO
FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                               DB 4;
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; LOCATION: (15)
; OTHER INFORMATION:
US-09-248-796A-20671
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PRIOR FILING DATE: 2000-02-01
NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CURAPAUSeqFormatter Version
SEQ ID NO 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20671, Ap
Patent No. 6747137
                                                                                                                                                             Matches
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 20671
LENGTH: 217
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                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/248,796A CURRENT FILING DATE: 1999-02-12 PRIOR APPLICATION NUMBER: US 60/074,725 PRIOR FILING DATE: 1998-02-13 PRIOR APPLICATION NUMBER: US 60/096,409 PRIOR APPLICATION NUMBER: US 60/096,409 PRIOR FILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Keith Weinstock et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 107196.132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/127,352
PRIOR FILING DATE: 1999-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 28208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number YML099C
                                                                                                                                                                                                                                                                                                                                                ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Saccharomyces
                                                                                                                                                                                                                                                                                                                             FEATURE:
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Local Similarity 22.7%;
                                                                                                                                                                                Local
                                      114 EVRLYKNKDMIQTLTNKSKIQNVGGT-YELHVGDTIDLIFNSEHQVLKNNTYWG 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        672 -ENIIGTDSLYGLPNSLILLFSDCVRIVRHNEYYNLTYLPVPR 713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      555 IFSFLKLIQDSTALDKVRAKEIVILPSEEDDNYKPLDTSNATTSSSEPRVDVVQEGLFRE
                                                                                 85 PKDWNDAEKQWEPPLIVNPKCATGCGPWEAPLIPNHDYI--GPWFPPDIKNPNYNGI--W
                                                                                                                       66 PSKWQMASS--EPP-----
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                                                                                                                                                                                Similarity
  TPRLIPNPDYYQVKTPGKLDKPIGGIGFEL-----
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                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09248796A
                                                                                                                                                                                                                                                             Identity of amino acid sequences at the
                                                                                                                                                                                8.1%;
27.2%;
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Pred. No. 6;
5; Mismatches
                                                                                                                                                                                Score 77; DB 4; Pred. No. 0.84;
                                                                                                                   -CVNKVSDWKLEILQNGLYLIYGQVAP----NANYNDVAPF
                                                                                                                                                             Mismatches
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                                                                                                                                                           Gaps
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RESULT 8
5395759-2
;Patent No. 5395759
; APPLICANT: HOLMES, IAN H.;DYALL-SMITH, MICHAEL L.
; APPLICANT: HOLMES, DNA SEQUENCES AND AMINO ACID SEQUENCE;
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Patent No. 6086880
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
INFORMATION FOR SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 326 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: ADLER, REID G.
REGISTRATION NUMBER: 30,988
REFERENCE/DOCKET NUMBER: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 22.8%; Pred. No. 1.6; hes 42; Conservative 29. Minarity
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FILING DATE: 07-JUL-1993
CLASSIFICATION: 530
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TOPOLOGY: unknown
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OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: MORRISON & FOERSTER STREET: 2000 Pennsylvania Avenue,
                                                                                                                                                                                                                174 QFIS 177
                                                                                                                                                                            168
                                                                                                                                                                                                                                                      130 NPQLYCDYNVV---
                                                                                                                                                                                                                                                                                                                                70 STSGEPFLTSTLCLYYPAEAKNEISDDEWENTLSQLFLTKGWPIGSVYFKDYNDINTFSV 129
                                                                                                                                                                                                                                                                                                                                                                                                         10 LTILISIILLNYILKTITNINDYIIFRELLLIALISPFVRTQNYGMYLPITGSLDAVYIN 69
                                                                                                                                                                                                                                                                                                                                                                                                                                               26 LWLFCSIVMLLFLC----SFSWLIFIFLQLETAKEPCMAK-----FGPLPSKWQM 71
                                                                                                                                                                                                                                                                                                                                                                     72 ASSEPP-----CV-----NKVSDWKLEILQNGLYLIYGQVAPNA---NYNDVAPFEV 115
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FRENCHICK, PATRICK J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         unknown
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                                                                                                                                                                                                                                                      -LMRYDNTSELDASELADLILNE--
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RESULT 9
US-08-496-944-2
                                  ; MOLECULE TYPE:
US-08-496-944-2
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 Query Match
                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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FILING DATE: 29-APR-1985
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 824,704
                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 14
CURRENT APPLICATION DATA:
                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 2763 amino acids
                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                               NAME: Elmer, James Scott
REGISTRATION NUMBER: 36,129
REFERENCE/DOCKET NUMBER: CG-
PRADATTON FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Hawthorne STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 22.8 nes 42; Conservative
                                                                         TOPOLOGY:
                                                                                            TYPE:
                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: CIBA-Geigy Corporation STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 04-FEB-1987
                                                                                                                                                                                                                                                             FILING DATE
                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10532
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26 LWLFCSIVMLLFLC----SFSWLIFIFLQLETAKEPCMAK-----FGPLPSKWQM 71
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 8.1%;
                                                                                                                                                                                                                                                                                US/08/496,944
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; Pred. No. 1.6;
29; Mismatches
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 Score 77;
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                                                                                                                                                                                                                                                                                                                     Version
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Length 2763;
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GENERAL INFORMATION:
                                                                                                                                                                                  Matches
                                                                                                                                                                                               Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ehrenshaft, Marilyn
APPLICANT: Jenns, Ann E.
TITLE OF INVENTION: Isolated Ge-
TITLE OF INVENTION: Resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Virginia C.
STREET: PO Box 37428
                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: BENDETT, VIRGINIA C.
REGISTRATION NUMBER: 37,092
REFERENCE/DOCKET NUMBER: 540
                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                          TYPE: 8
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CITY: Raleigh
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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   297
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                                                                                                                                                                                                                                                                                     amino acid
 QRLRSPLIMFWEHVA-
                           --LQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLTNKSKI------
                                                         NADSDRWLAQYIRAERLCEEVSEQVDLTNTSFYRDVADPATRNPVQTCRNKILNWKMGVP 296
                                                                                                                    SYESVEG-----WRAWLGCHVLSVSMAIFMRKSMTASWT----EQHEQAR--LMLQYSPL
                                                                                                                                                 SHSRTQGAQRSSWKLWLFC---SIVMLLFL---CSFSWLIFIFLQLETAKEFCWAKFGFL
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                                                                                         ---PSKW--
                                                                                                                                                                                                                                                                                                        550 amino acids
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Ehrenshaft, Maril
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Pred. No. 5.1;
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TAYMHEPVLHTATNKDSFTAPYLAERLSLTDF
                                                                                                                                                                                                            DB 3;
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                                                                                       --QMASSEP--PCVNKVSDWKLEI-
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                                                                                                                                                                                                             Length 550;
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US-09-328-352-5760
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Best Local S
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APPLICANT: GATY L. BECTON et al.
APPLICANT: GATY L. BECTON et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 5760
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Patent No. 6562958
                                                                                                                                 Best
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                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                              SEQ ID NO 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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CURRENT FILING DATE: 1999-11-18
PRIOR APPLICATION NUMBER: 60/109,283
PRIOR FILING DATE: 1998-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Famodu, Omolayo O.
APPLICANT: Orozco, Buddy
APPLICANT: Rafalski, Antoni
APPLICANT: Shen, Jennie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                   SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Sterol Metabolism Enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                    ORGANISM: Nicotiana tabacum
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                                                                                                                                                                                                                                         LENGTH: 2'
TYPE: PRT
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                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   131 FIYILRN-LLLTYWQVSLYVGCTVSWIAFLYYAFYVNQKDIKIFSKKFK-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80 SIVLFLFGISLMPVTYIILNI-----VNKY--LPSHEKIRDDLPFPKISIITFIASIW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31 SIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSKWQMASSEP-PCVNKV----SDW 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42;
                                                                                                                                                                                                                                                              271
                                                                                                                    h 7.8%; Score 74.5; DI Similarity 22.7%; Pred. No. 2.3; 34; Conservative 23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                       PLPH-MLQG-----WLRNYIGGVLLYFISGFLWCFYIYHLKRNVYIPKDAIPSNKAM 81
                                                                             PISHSRTQGAQRSSWKLWI--FCSIVMLLFLCSFSWLIFIF-----
-LQLETAKEPCMAKFGPLP--SKWQMASSEPPCVNKVSD--WKLEILQNGLYLI---YGQ 100
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Pred. No.
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                                                                                                                      43;
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RESULT 14
US-08-349-357-2
; Sequence 2, Application US/08349357
; Patent No. 6265556
; Patent No. 6265556
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/349,357
FILLING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1203
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linco
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CLASSIFICATION: 530
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/697,610
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ADDRESSEE: Campbell and Flores
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APPLICANT: Sato, Takaaki
TITLE OF INVENTION: CD40 Associated Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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o. 6172187
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Pred. No. 6.6;
20; Mismatches
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RESULT 15
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GENERAL INFORMATION:
APPLICANT: Wisnieski, Bernadine J
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Best Local
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TELEPAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/349,35:
FILING DATE: 02-DEC-1994
CLASSIFICATION: 435
                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1203
TELECOMMUNICATION INFORMATION:
COMPUTER: VE System 386
OPERATING SYSTEM: MS-DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Floppy disk
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ZIP: 92122
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California
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Pred. No. 6.6;
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CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/852,625

FILING DATE: 12 MARCH 1992

ATTORNEY/AGENT INFORMATION:

NAME: Donald G. Lewis

REGISTRATION NUMBER: 28636

REFERENCE/DOCKET NUMBER: BJW-2

TELECOMMUNICATION INFORMATION:

TELEPHONE: (619) 554-2421

TELEPHONE: (619) 554-2421

TELEPAX: (619) 554-6312

INFORMATION FOR SEQ ID NO: 6:

SEQUENCE CHARACTERISTICS:
LENGTH: 157 amino acids

TYPE: AMINO ACIDS

TOPOLOGY: linear

MOLECULE TYPE: protein

FRATURE:
                                                                                                                                                                                                                                                                                                                                 Query Match 7.8%; Score 74; DB 5; Length 157; Best Local Similarity 24.4%; Pred. No. 1.2; Matches 30; Conservative 21; Mismatches 32; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Tumor Necrosis Factor (feline)
PUBLICATION INFORMATION:
AUTHORS: McGraw, R. A., Coffee, B.W., Otto,
AUTHORS: C.M., Drews, R.T. and Rawling, C.A.
TITLE: Gene Sequence of Feline Tumor Necrosis
TITLE: Factor '.
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PAGES: 5564
DATE: 190
RELEVANT RESIDUES IN SEQ ID NO: 6: 1-157
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TNF5_CHICK
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AA608050
Q861B8
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AAH70042
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Q65605
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Q918d8 gallus gall
Q741w5 lactobacill
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SEQUENCE SEQUENCE SEQUENCE SEQUENCE Clark IN Chen J. Eaton D. Huang Lewis L. Seshagi Vandles Seshagi Vandles Sefort Si S., Godowsk The see effort bioinfo Genome [3] SEQUENCE	SEQUITISSI MEDL: Gurne Balde Balde Godda "Ides and : Curr	.∞ ⊂	Ο C C C C C C C C A A A A A A A A A A A
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Med=12975309; L., Abaya E., L., Crowley Grimaldi C., Climowski L., Climowski L., Singh J. C., Wieand D. C., Wieand D. Zhang M., Z discovery in discovery in discovery in movel human sec ment.", 1270(2003). MM N.A. MM N.A. Med=10037686; J., Yu GL. Cwon B.S., Tovel activat tor superfami 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999) 156-6061(1999)	i=10074428, A., Huang Dowd P., Baker K.! member of nn ortholog	QD; Created) Last sec Last and Ligand si gland) (h 19/PRO175 3=AITRL, nordata; cimates;	00400000000000000000000000000000000000
wley C wley C C. Gu C. Gu bbie E h J., Zhai y init y entity y entity family 999) cated T atted T	10074428; ., Huang A., P Dowd P., Brush Baker K.P., Go member of the ortholog of m	PRT; uence otatic otatic uperfa GITRL) GITRL, Catarr	Q72VG7 Q8F914 AAS68957 Q805N8 Q80AT3 PROB OCEIH Q8E0H3 Q8E0H3 Q8EDB7 P89064 Q8IDP7 P89064 Q8IDP7 P89064 QBIDP7 P89164 QBIDP7 P89164 QBIDP7 P89164 QBIDP7
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GO; GO:0007267; P:cell-cell signaling; TAS.
GO; GO:0007165; P:signal transduction; TAS.
InterPro; IPR006052; TWF family.
InterPro; IPR008983; TNF_like.
PROSITE; PS00251; TWF_1; FALSE_NEG.
PROSITE; PS00049; TWF_2; FALSE_NEG.
Cytokine; Glycoprotein; Signal-anchor; Transme DOMAIN
1 28 Signal-anchor for TRANSMEM 29 49 Signal-anchor for
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24-MAY-2004 (TrEN
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TNUESF18 protein
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CARBOHYD
SEQUENCE
TISSUE-Synthetic constructs;
MEDLINE-22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse I
Klausner R.D., Collins F.S., Wagner L.,
Altschul S.F., Zeeberg B., Buetow K.H.,
                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- SUBCELLULAR LOCATION: Type II membrane protein (Potential).
-!- TISSUB SPECIFICITY: Expressed at high levels in the small intestine, ovary, testis, kidney and endothelial cells.
-!- INDUCTION: Up-regulated after stimulation by lipopolysaccha-
-!- SIMILARITY: Belongs to the tumor necrosis factor family.
                                                                                   SEQUENCE
                                                                                                                                                                  Homo sapiens (Human)
                                                                                                                                                                                  INFSF18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AF125303; AAD22634.1; -.
EMBL; AY358868; AAQ89227.1; -.
EMBL; AF117713; AAD19695.1; -.
Genew; HGNC:11932; TNFSF18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MCLSHLENMPLSHSRTQGAQRSSWKLWLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMA
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Signal-anchor for type II membrane protein (Potential).

Extracellular (Potential).

N-linked (GlcNAc. . .) (Potential).

N-linked (GlcNAc. . .) (Potential).
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Pred. No. 1.2e-80;
                                                                                                                                  Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3D78CE6B90F4C9E3 CRC64;
Grouse L.H., Derge J.G.,
gner L., Shenmen C.M., Schuler G.D
ow K.H., Schaefer C.F., Bhat N.K.,
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RC TISSUE=PCR rescued clones;

RMEDIINE=22388257; pubMed=12477932;

RMEDIINE=22388257; pubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Altschul S.F., Zeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Brapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M.G., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RB Brownstein M.G., Usdin T.B., Foshiyuki S., Carninci P., Prange C.,

RB Brownstein M.G., Usdin T.B., Foshiyuki S., Carninci P., Prange C.,

RB Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Worley K.C., Hale S., Garcia A.M., Gab L.J., Hullyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gibbs R.A.,

RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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01-JUN-2004 (TrEMBLrel. 27, Last sequence update)
01-JUN-2004 (TrEMBLrel. 27, Last annotation update)
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"Generation and initial analysis of
"Generation cDNA sequences.";
proc. Natl. Acad. Sci. U.S.A. 99:16
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Pred. No. 1.
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                                             EMBL; AY234223; AAO89011.1; -.
EMBL; AJ577579; CAB12166.1; -.
EMBL; AJ577580; CAB12167.1; -.
MGD; MGI:2673064; Thifsf18.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0015164; F:tumor necrosis fac
GO; GO:0006955; P:immune response; II
InterPro; IPR008983; TNF_like.
InterPro; IPR003636; TNF_subf.
                                                                                                                                                                                                                                                                                                                                     Q80YG2;
01-JUN-2003
01-JUN-2003
05-JUL-2004
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Bianchini R., Nocent
Submitted (FEB-2003)
                                                                                                                                                                                                                                                                                      Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                      Proc. Natl. Acad. Sci.
                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                   Name=Tnfsf18; Synonyms=tnfsf
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z J., Myers R.M.,
s D.E., Schnerch
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Murinae; Mus
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Proc. Natl. Acad. Sci. U.S.A.
EMBL; AJ577579; CAE12166.1; -
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Mammalia; Eutheria;
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SEQUENCE FROM
STRAIN=CBA/Ca;
                                    NCBI_TaxID=10090;
                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Mouse glucocorticoid-induced tumor
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Rodentia;
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Pred. No. 5.6e-31;
0; Mismatches 58;
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Sciurognathi; Muridae;
                                                                          Craniata; Vertebrata; Sciurognathi; Muridae;
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STRAIN-BALL, ...

PubMed=14647196;

A Kim J.D., Choi B.K., Bae J.S., Lee U.H., ....

A Youn B.S., Vinay D.S., Kwon B.S.;

"Cloning and characterization of GITR ligand RL Genes Immun. 4:564-569(2003).

DR EMBL; AY267900; AAP96745.1; ...

DR GO; GO:0016020; C:membrane; IEA.

GO; GO:0015164; F:tumor necrosis factor rec GC; GO:0005955; P:Immune response; IEA.

DR InterPro; ITRR00336; TNF subf; 1.

DR ProDom; PD002012; TNF subf; 1.
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Q7TNY2;
01-OCT-2003
01-OCT-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                         GITR ligand.
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Waldmann H.;

"Mouse glucocorticoid-induced tumor necrosis factor receptor ligand costimulatory for T cells.";
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EMBL; AJ577580; CAE12167.1; -.
Alternative splicing.
SEQUENCE 173 AA; 19745 MW; 1FD2294CACF424D2 CRC
                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
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VLQTLMNDFQILPIGGVYELHAGDNIYLKFNSKDHIQKNNTYWGIILMPDLPFIS 173
                  MIQTITIKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 177
                                              SSSKWHMTSPKPHCVNTTSDGKLKILQSGTYLIYGQVIPVDKKYIKDNAPFVVQIYKKND
                                                            LPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAP-NANY-NDVAPFEVRLYKNKD
                                                                                             MEEMPLRESSPORAERCK-KSWLLCIVALLLMLLCSLGTLIYTSLK-PTVIESCMVKPEL
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Rodentia;
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Pred. No. 1.3e-30;
0; Mismatches 59
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Sciurognathi; Muridae;
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Submitted (AUG-2003) to the EMBL/GenBank/
EMBL; AY320040; AAP70494.1; -.
EMBL; AY320040; AAP55265.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0005164; F:tumor necrosis factor ro
GO; GO:000515; P:immune response; IEA.
InterPro; IPR003636; TNF subf.
ProDom; PD002012; TNF_subf; 1.
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Q7TS55;
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02-MAR-2004 (TrEMBLrel. 27, C
02-MAR-2004 (TrEMBLrel. 27, I
02-MAR-2004 (TrEMBLrel. 27, I
Glucocorticoid-induced-tumor
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Submitted
                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
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Eukaryota; Metazoa;
Mammalia; Eutheria;
   SEQUENCE FROM N.A.
                                                    NCBI_TaxID=10090;
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(TYEMBLIEL. 25, Last sequence update)
(TYEMBLIEL. 27, Last annotation update)
(Glucocorticoid-induced-tumor necrosis factor receptor
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or necrosis factor
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Sciurognathi; Muridae
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A Tregaskes C.A., Young J.R., Burnside J.;
Tregaskes C.A., Young J.R., Burnside J.;
U. Submitted (MAR-2003) to the EMBI/GenBank/DDBJ databases.
C --- FUNCTION: Cytokine that binds to TNFRSF5. Mediates B-cell production in the presence of co-stimulus as well as IgE production in the presence of IL-4. Involved in immunoglobulin C class switching (By similarity).
C --- SUBGUNIT: Homotrimer (By similarity).
C --- SUBGUNIT: Homotrimer (By similarity).
C --- SUBGUNIT: Homotrimer (By similarity).
C --- SUBGUNIT: Soluble form (By similarity).
C --- SUBGUNIT: Soluble form (By similarity).
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MEDLINE=22883945; PubMed=14521928;
Yu K.Y., Kim H.S., Song S.Y., Min S.S., Jeong J.J., Youn B.S.;
Yu Kern, Kim H.S., song S.Y., Min S.S., Jeong J.J., Youn B.S.;
"Identification of a ligand for glucocorticoid-induced tumor necrosis factor receptor constitutively expressed in dendritic cells.";
Biochem. Biophys. Res. Commun. 310:433-438(2003).
                                           use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2003 (Rel. 41, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Tumor necrosis factor ligand superfamily memb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L) (CD154 protein).
Name=TNFSF5; Synonyms=CD40LG,
Gallus gallus (Chicken).
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                                                                                                                                                     s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the En European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                      proteolytic processing (By similarity). SIMILARITY: Belongs to the tumor necrosis factor family.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CD40 ligand)
                                                                                                                                                                                                                                                                                                                                                                                                                                         immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                             exists
                                                                                                                                                                                a collaboration
                                                                                                    for
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                                                                           .ch/announce/
                                                                                                       commercial
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RESULT 11
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Best Local S
Matches 36
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GO; GO:0016021; C:integral to membrane; ISS.
GO; GO:0005174; F:CD40 receptor binding; ISS.
GO; GO:0042100; P:B-cell proliferation; ISS.
GO; GO:0042100; P:B-cell proliferation; ISS.
GO; GO:0006954; P:inflammatory response; ISS.
GO; GO:0007159; P:leukocyte cell adhesion; ISS.
GO; GO:0007169; P:leukocyte cell adhesion; ISS.
GO; GO:0030168; P:platelet activation; ISS.
InterPro; IPR003263; TNF_5.
InterPro; IPR003263; TNF_family.
InterPro; IPR003636; TNF_subf.
Pfam; PF00229; TNF; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00229; INF; ...

PRINTS; PR01702; CD40LIGAND.

PRODOM; PD008600; TNF 5; 1.

ProDom; PD002012; TNF subf; 1.

PROSITE; P800251; TNF 1; 1.

PROSITE; P800251; TNF 2; 1.

PROSITE; P850049; TNF 2; 1.

PROSITE; PS50049; TNF 2; 1.

Cytokine; Glycoprotein; Signal-anchor; Transmembrane.

Cytokine; Glycoprotein; Signal-anchor; Transmembrane.

Cytokine; Glycoprotein; Signal-anchor; Transmembrane.
                                                                                                                                                                                                   Q74LW5
Q74LW5;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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CARBOHYD
CARBOHYD
CARBOHYD
SEQUENCE
       Pridmore R.D., Berger B., Desiere F., Vilanova D., Pittet A.-C., Zwahlen M.-C., Rouvet M., Altermann Mollet B., Mercenier A., Klaenhammer T., Arigoni F. The genome sequence of the probiotic intestinal F. Lactobacillus johnsonii NCC 533.";
                                                                                                                                                                        Hypothetical protein.
OrderedLocusNames=LJ0068;
                                                                     STRAIN=NCC 533;
PubMed=14966310;
                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DOMAIN
SITE
                                                                                                                                                               Lactobacillus johnsonii.
                                                                                                                          NCBI_TaxID=33959;
                                                                                                                                                 Bacteria; Firmicutes;
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                                                                                                                                                                                                                                                                                                                                                                                                                    KW-MTTSYAPTSSLISYHEGKLKVEKAGLYYIYSQVSFCTKAAASAPFTLYIYLYDMEE
                                                                                                                                                                                                                                                                                                                                                                                                                                               KWQMASSEPPCVNKVS--DWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLY-----
Acad.
                                                                                                                                                                                                   (TrEMBLrel.
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110
190
124
146
251
                                                                                                                                                                                                                          (TrEMBLrel.
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229
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U.S.A.
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27,
27,
                                                                                                                                                                                                 Last sequence update)
Last annotation update)
                                                                                                                                                                                                                              Created)
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Pred.
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Extracellular (Potential).

Cleavage (By similarity).

Potential.

N-linked (GlcNac. . .) (Potential).
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Signal-anchor for type II membrane
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumor necrosis factor ligand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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1 (GlcNAc...)
1 (GlcNAc...)
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                                                           Vilanova D.,
                                                                                                                                                   Lactobacillaceae
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                                  n E., Barra
F., Schell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                           Barretto
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RESULT 12
AASS080
ID AASS0
AC AASS0
AC AASS0
DT 02-M
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RESULT 13
Q86IB8
ID Q86IB
AC Q86IB
DT 01-JU
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DT 01-MA
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Best Local S
Matches 42
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                     Q86IB8;
01-JUN-2003
01-JUN-2003
01-MAR-2004
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02-MAR-2004
02-MAR-2004
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Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pridmore R.D., Berger B., Desiere F., Vilanova D., Barretto C., Pittet A.-C., Zwahlen M.-C., Rouvet M., Altermann E., Barrangou R., Mollet B., Mercenier A., Klaenhammer T., Arigoni F., Schell M.A.; "The genome sequence of the probiotic intestinal bacterium Lactobacillus johnsonii NCC 533."; Lactobacillus johnsonii NCC 533."; Proc. Natl. Acad. Sci. U.S.A. 101:2512-2517(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hypothetical protein. SEQUENCE 273 AA; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=14966310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Firmicutes; Lactobacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS08050;
  Hypothetical
                                                                                                                  Q86IB8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=33959;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lactobacillus johnsonii.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WIF----LIVFIG-----INIFLGIELWQTPTLLSAGSTETQTDIKSETSADQTTIPKVN 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proteome; Hypothetical protein.
273 AA; 31108 MW; 34414AF96D1747A8 CRC64;
                                                                                                                                                                                                                                  REVMRFKDDSQNVYQGKNYTYLSELSEGD--DYIFNQKTKYGEFFAATARLHIIVKDN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DKQ----EDGYYLAAKNDDSWIKKATQQVKGQVENNSSENLISVHLDKPITLSKNKKEAL 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DWKLEILQNGLYL---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             WLFCSIVMLLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSKWQM---ASSEPPCVNKVS
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                       (TrEMBLrel.
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                                                                  (TrEMBLrel.
  protein.
                                                                                                                    PRELIMINARY,
                                                                                                                                                                                                                                                                                                                             EDGYYLAAKNDDSWIKKATQQVKGQVENNSSENLISVHLDKPITLSKNKKEAL 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31108 MW;
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24,
26,
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27,
27,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 88.5; DB Pred. No. 5.9; 32; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Created)
Last sequence update)
Last annotation updat
                       Last sequence up
                                                                  Created)
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Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   34414AF96D1747A8 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                    516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lactobacillaceae;
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                                             update)
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                                                                                                                                                                                                                                                                               ----HQVLKNN 162
                                                                                                                                                                                                                                                                                                                                                                          -PFEVRLYKNKDM 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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ID ACCOMENTATION OF THE PROPERTY OF THE PROPER
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Best Local :
                                                                                                                                                                                                                                                            Matches
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Best Local
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01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Kila protein, putative phage-related DNA binding pr
Name=kila; OrderedLocusNames=CTC01071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q896D6;
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Gloeckner G., Eichinger L., Szafranski K., Pachebat J., Dear P.,
Lehmann R., Baumgart C., Parra G., April J.F., Guigo R., Kumpf K.,
Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noegel A.A.
"Sequence and analysis of chromosome 2 of Dictyostelium discoideum.";
Nature 418:79-85(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases EMBL; ACI1655; AA052204.1; -. Hypothetical protein.
SEQUENCE 516 AA; 58667 MW; 80B308CF6CDF97C2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
NCBI_TaxID=44689;
[1]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=Massachusetts / E88;
MEDLINE=22457253; PubMed=12552129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=AX4;
                                                                                                                                                                                                                                                                                                                                                                                                     Complete proteome. SEQUENCE 270 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AE015939; AA035654.1;
InterPro; IPR003497; BRO_N.
Pfam; PF02498; Bro-N; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Proc. Natl. Acad.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=1513;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clostridium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clostridium tetani.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Decker I., Her
Gottschalk G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The genome sequence of Clostridium tetani, the causative agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 419
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                                                                                                       VNNNINEKNOFIIN----IPKDGNCDGIILLIN
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                                                                                                                                                                               KVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLTNKSKIQNVGGT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               n H., Baeumer S., Fricke W.F., W
Herzberg C., Martinez-Arias R.,
                                                                                                                                                                                                                                                                                                                                                                                                         270 AA; 31533 MW;
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25.4%;
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                                                                                                                                                                                                                                                            Score 85.5; D
Pred. No. 11;
18; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 19;
7; Mismatches
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YELHVG-----DTIDLIFNSEHQVLKNNTYW---GIILL
                                                                                                                                                                                                                                                                                                                                                                                                         C378AB524C0133A8 CRC64;
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                                                                                                                                                                                                                                                                36;
                                                                                                                                                                                                                                                                                                                                Length 270;
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, Henne A.,
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A., Noegel A.A.;
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RA Dunham I., Hunt A.R., Collins J.E., Brusklewich R., Beare D.M.,
RA Clamp M., Smink L.J., Almscough R., Almeida J.P., Babbage A.K.,
RA Bagguley C., Bailey J., Barlow K.F., Bates K.N., Beasley O.P.,
RA Bagguley C., Bailey J., Barlow K.F., Bates K.N., Beasley O.P.,
RA Burrill W.D., Burton J., Carder C., Carter N.P., Chen Y., Clark G.,
RA Burrill W.D., Burton J., Carder C., Carter N.P., Chen Y., Clark G.,
RA Clegg S.M., Cobley V.E., Cole C.G., Collier R.E., Comnor R.,
Conroy D., Corby N.R., Coville G.J., Cox A.V., Davis J., Dawson E.,
RA Conroy D., Corby N.R., Coville G.J., Cox A.V., Davis J., Dawson E.,
RA Dhami P.D., Dockree C., Dodsworth S.J., Durbin R.M., Ellington A.G.,
RA Evans K.L., Fey J.M., Fleming K., French L., Garner A.A.,
RA Evans K.L., Fey J.M., Fleming K., French L., Garner A.A.,
RA Hunt S.E., Jones M.C., Kershaw J., Kimberley A.M., King A.,
Laird G.K., Langford C.F., Leversha M.A., Lloyd C., Lloyd D.M.,
RA Hail R.E., Hall-Tamlyn G., Heathcott R.W., Ho, S., Holmes S.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA McClay J., McJaren S., McMurray A.A., Milne S.A., Mortimore B.J.,
RA Odell C.N., Pavitt R., Pearce A.V., Pearson D., Phillimore B.J.C.T.,
RA Phillips S.H., Plumb R.W., Ramsay H., Ramsey Y., Rogers L., Ross M.T.,
RA Scott C.E., Sehra H.K., Skuce C.D., Smalley S., Smith M.L.,
Soderlund C., Spragon L., Steward C.A., Sulston J.E., Swann R.M.,
RA Vaudin M., Wallis J.M., Whiteley D.L.,
RA Williams L., Williams S.A., Williamson H., Wilmer T.E., Wilming L.,
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Nomura N., Miyajima N., Sazuka T., Tanaka A., Kawarabayasi Y.,
Sato S., Nagase T., Seki N., Ishikawa K.-I., Tabata S.;
"Prediction of the coding sequences of unidentified human genes.
The coding sequences of 40 new genes (KIAA0001-KIAA0040) deduced
analysis of randomly sampled cDNA clones from human immature myel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1998 (Rel. 36, Created)
05-JUL-2004 (Rel. 44, Last sequence up
05-JUL-2004 (Rel. 44, Last annotation
Membrane protein MLC1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=2122590; PubMed=11326298; DOT=10.1038/sj/mp/4000869; Meyer J., Huberth A., Ortega G., Syagailo Y.V., Jatzke S., Moessner R., Strom T.M., Ulzheimer-Teuber I., Stoeber G., Sch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pedigree.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         channel is associated
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15-JUL-1998 (Re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Hippocampus;
MEDLINE=21225990; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Construction of expression-ready cDNA curation of 330 KIAA cDNA clones.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nakajima D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=MLC1; Synonyms=WKL1, KIAA0027; Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Bone marrow;
MEDLINE=96051387; P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of 330 KIAA cDNI
9:99-106(2002).
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Q96RP5; Q9UGY8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6:302-306(2001)
                                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=10591208; DOI=10.1038/990031; R., Collins J.E., Bruskiewich R., Be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE SPECIFICITY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       catatonic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    encoding a putative cation schizophrenia in a large
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A clones for
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or KIAA
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RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Wullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Garcia B.D., Dickson M.C.,
RA Rathey J., Hulton E., Ketteman M., Sodergran E.J., Lu, K., Schall R., Scheck S., M., Krzywinski M.I., Skalska U., Smail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RA Minoshima S., Kawasaki K., Sasaki T., Abakawa S., Kudoh J.,
RA Minoshima S., Kawasaki K., Sasaki Y., Abki N., Mitsuyama S.,
RA Roe B.A., Chen F., Chu L., Crabtree J., Deschamps S., Do A., Do T.,
RA Dorman A., Fang F., Fu Y., Hu P., Hua A., Kenton S., Lai H., Lao H.I.,
RA Lewis J., Lewis S., Lin S.-P., Loh P., Malaj E., Nguyen T., Pan H.,
RA Phan S., Qi S., Qian Y., Ray L., Ren Q., Shaull S., Sloan D., Song L.,
RA Wang Q., Wang Y., Wang Z., White J., Willingham D., Wu H., Yao Z.,
RA Fulton R., Johnson D., Bemis G., Bentley D., Bradshaw H., Bourne S.,
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RA Hinds K., Kemp K., Latreille P., Layman D., Ozersky P., Rohlfing T.,
RA Scheet P., Walker C., Wamsley A., Wohldmann P., Pepin K., Nelson J.,
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RA Chan A.S., Lane L., Tilahun Y., Wright H.;
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MEDLINE-2115271; Pubmed-11254442;
Leegwater P.A.J., Yuan B.Q., van der Steen J., Mulders
Koenst A.A.M., Ilja Boor P.K., Mejaski-Bosnjak V.,
van der Maarel S.M., Frants R.R., Oudejans C.B.M., Sch
"Identification of novel mutations in MLC1 responsible for megalencephalic leukoencephalopathy with subcortical cysts.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=22608298; PubMed=12665801; DOI=10.1038/nbt810; Gevaert K., Goethals M., Martens L., Van Damme J., StaThomas G.R., Vandekerckhove J.;
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[5]
                                                                                                van der Knaap M.S.;
                                                                                                                                VARIANTS MLC SER-91; LYS-140 AND SER-140.
PubMed=11935341; DOI=10.1007/s00439-002-0682-x;
Puegwater P.A.J., Boor P.K.I., Yuan B.Q., van des
Konst A.A.M., Oudejans C.B.M., Schutgens R.B.H.,
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                                                                                                                                                                                                                                                                                                                                              Pronk J.C., van der Knapp M.S.;
"Mutations of MLC1 (KIAA0027), encoding a putative membrane protein, sause megalencephalic leukoencephalopathy with subcortical cysts.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Exploring proteomes and analyzing pro
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                                                                                                                                         Pronk
                                                                                                                                    Steen J.,
Pronk J.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptides.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schutgens
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                                                                                                                                                                                       Visser
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VARIANT MLC GLU-58

Genet. 110:279-283 (2002).

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C:- TISSUE SPECIFICITY: Expressed in the brain, with highest levels found in the amygdala, nucleus caudatus, thalamus and hippocampus.

C:- DISEASE: Defects in MLC1 are a cause of megalencephalic leukoencephalopathy with subcortical cysts (MLC) [MIM:604004]. MLC is an autosomal recessive disorder characterized by macrocephaly, deterioration of motor functions with ataxia, and spasticity, eventuating in mental decline. The brain appears swollen on magnetic resonance imaging, with diffuse white-matter abnormalities and the invariable presence of subcortical cysts.

C:- DISEASE: Defects in MLC1 are a cause of periodic catatonia (PC), a familial subtype of catatonic schizophrenia which is a genetically heterogeneous disorder characterized by psychosis and psychomotor disturbances.
                                                                                                        Query Match
Best Local Similarity
                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an entitles requires a license agreement (See http://www.isb-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Megalencephalic leukoencephalopathy with subcortical cysts; a founder effect in Israeli patients and a higher than expected carrier rate among Libyan Jews.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=12189496; DOI=10.1007/s00439-002-0770-Y;
Ben-Zeev B., Levy-Nissenbaum E., Lahat H., Anikster Y., Shinar Y.,
Brand N., Gross-Tzur V., MacGregor D., Sidi R., Kleta R., Frydman M.,
Pras E.;
                                                                                                                                                                                                                                                                                                                                                                              INIT_MET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation between Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                    VARIANT
                                                                                                                                                                                                VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                          GO; GO:0000299; C:integral Direct protein sequencing; Transmembrane; Transport.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        MIM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; BC028425;
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EMBL; AF319633; AAK60119.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- FUNCTION: May be a transporter. May act as a non-selective
neuronal cation channel.
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AL022327;
                               126 IWFGCKLVLNPSAININFNLILLLLELLMAATVIIA-----ARSSEEDCKKK-- 173
 84 DWKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLTNKSKIQNVGGTYELH 143
                                                           26 LWLFCSIVM--LLFLCSFSWLIFIFLQLETAKEPCMAKFGPLPSKWQMASSEPPCVNKVS 83
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112
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149
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58
                                                                                          Conservative
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CAB51559.1; ALT_INIT.
AAH28425.1; --
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p -> S (in MLC).
/FTId=VAR_017439.
S -> L (in MLC).
/FTId=VAR_011699.
T -> R (in MLC).
                                                                                                          Score 85.5;
Pred. No. 16;
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                                                                                                                        DB 1;
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DЬ	Ϋ́	Db
218 VDDSVSGPHLSVTEFWILVACEPSAIA 244	144 VGDTIDLIENSEHQVLKNNTYWGIILLANPQFIS 177	

Search completed: November Job time: 196 secs 'n 2004, 12:40:57

GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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A_Geneseq_23Sep04:*
1: geneseqp1990s:*
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1 ETAKEPCMAKFGPLPSKWQM.....
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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75 100.0	169	N		man e
100.	169	ω	AAB08785	A hum
100.	169	w	AAY53061	Human en
100.	169	σ	ADA11464	Human
100.	169	7	ADC01858	Human
100.	177	N	AAY15817	Aay15817 Amino aci
100.	177	Ŋ	AAY06646	Human
100.	177	4	AAB47056	Aab47056 PRO175. 5
100.	177	4	AAB20109	
100.	177	4.	AAB53065	Aab53065 Human ang
100.	177	4	AAB47287	Aab47287 PRO175 po
100.	177	4	AAB50979	Human E
100.	177	v	AAU81953	Aau81953 Human PRO
100.	177	თ	ABR42323	Human
100.	177	9	ABP60544	Abp60544 Human tum
100.	177	7	ADC35222	
100.	177	7	ABW02285	Abw02285 Human end
83.	508	4	ABG17426	Novel
12.	1221	7	ADC94332	
11.	599	σ	ABP25933	Str
11.	183	ω	AAG43533	Aag43533 Arabidops
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KW autoimmune disease; lupus; multiple sclerosis; chronic inflammation;

KW acute inflammation; acute allograft rejection; graft versus host disease;

KW transplant rejection; foetal resorption; faecal peritonitis; allergy;

KW bowel disease; sepsis; leukaemia; chronic hypergammaglobulinemeia;

KW polychondritis; scleroderma; Wegener granulomatosis; dermatomyositis;

KW chronic active hepatitis; myasthenia gravis; psoriasis; vitilogo;

KW Steven-Johnson syndrome; idiopathic sprue; gluten-sensitive enteropathy;

KW gemphigus vulgaris; Goodpasture's disease; bullous pemphigoid;

KW discoid lupus; dense deposit disease; endocrine ophthalmopathy;

KW wirritable bowel disease; asthma; Grave's disease; sarcoidosis; cirrhosis;

KW lymphopenias; polyarteritis nodosa; Sjogren's syndrome; Hechet's disease;

KW primary myxedema; polymyositis; mixed connective tissue disease;

KW keratoconjunctivitis sicca; vernal keratoconjunctivitis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yu G,
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                                                                        Homo sapiens
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Pred. No. 2.2e-69;
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Location/Qualifiers

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AAY53061 standard; protein; 169 AA

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                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a human endokine-alpha polypeptide. The polypeptide is a cytokine which is similar to tumour necrosis factor (TNP). The endokine-alpha polypucleotides and polypeptides are useful for treating AIDS, chronic lymphocyte disorder, tumours, parasitic disease, treating AIDS, chronic lymphocyte disorder, tumours, parasitic disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polynucleotide encoding endokine alpha protein, which is a member of tumor necrosis factor useful for treating inflammatory diseases, disorders such as cancer, allergy, diabetes and various neurological
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28-MAY-1999;
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                                                                                                                                                                                                                                                                                      Sequence 169 AA;
                                                                                                                                                                                                                                                                                                                          keratoconjunctivitis,
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164
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                                  NPQFIS 126
                                                                    PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                            PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
                                                                                                                                                              ETAKEPCMAKFGPLPSKWQMASSBPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA 60
                                                                                                                                            ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                   Conservative
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99US-0136788P
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                                                                                                                                                                                                                100.0%; Score 675; DB 3;
100.0%; Pred. No. 2.2e-69;
tive 0; Mismatches 0;
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AAY53061;

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RESULT 4
ADAL146
ID ADAL1
AC ADAL1
AC ADAL1
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AC ADAL1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents human endokine alpha which is a member of the tumour necrosis factor (TNF) family of cytokines. Endokine alpha protein and polynucleotides can be used in diagnostic and therapeutic methods concerning TNF family-related disorders. These include disorders associated with immunomodulation and inflammation, cell proliferation, and concerning the immunomodulation and inflammation.
                human; endokine alpha; melanoma; sarcoma; tumour; tumour regression; infection; viral; bacterial; yeast; fungal; Toxoplasma gondii; Schistosoma manson; Listeria monocytogenes; tumour necrosis factor; TNF-related disorder; endokine alpha-related disorder; immunomodulation;
                                                                                                                                                                                                                06-NOV-2003
                                                                                                                                                                                                                                                                                                                     ADA11464 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 169 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               angiogenesis, tumour metastasis, apoptosis, sepsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated human endokine alpha
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                                                                                                                                                          endokine alpha.
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                                                                                                                                                                                                                                                                                                                  protein; 169
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Pred. No. 2.2e-69;
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Matches 126; Conserv

Conservative

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Mismatches

100.0%; Score 675; DB 6; 100.0%; Pred. No. 2.2e-69;

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ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA ETAKEPCWAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA

Sequence

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The invention relates to an isolated human endokine alpha polypeptide. A CC cell recombinant for the human endokine alpha polypeptide is useful for producing the protein by recombinant techniques. The antigenic epitope producing the protein by recombinant techniques. The antibodies concluding memoral polypeptides are useful to raise antibodies. CI including memoral antibodies. The peptides and antipeptide antibodies are used in a variety of qualitative or quantitative assays for the protein. The protein is useful for tumour targeting and thus used in patients with melanoma and sarcoma for tumour regression and extension of patients with melanoma and sarcoma for tumour regression and extension of comparison the protein is also useful for treating viral, bacterial, yeast, fungal and other infections e.g., The protein is also useful for treating other tumour necrosis factor (TNF)-related disorders. The nucleic acid is useful as probes for gene mapping by in situ hybridisation and for detecting expression of endokine alpha related disorder such as also useful for diagnosing an endokine alpha-related disorder such as also useful immunomodulation and inflammation, cell of turnoling them to the protein is also cald is also useful for diagnosing an endokine alpha-related disorder such as also useful for diagnosing an endokine alpha-related disorder such as also useful and compared with immunomodulation and inflammation, cell of the protein associated with immunomodulation and inflammation.
                                                                            proliferation, angiogenesis, tumour metastasis, apoptosis, sepsis or endotoxaemia. The antibody is useful diagnostically or therapeutically as antagonists in the treatment of alpha and/or TNF-related disorders. The antibody is also useful for purification of the protein. The antibodies are useful for detecting the protein and for tracking the fate of various regions of a protein precursor which undergoes post-translational
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated endokine alpha polypeptide, a member of tumor necrosis factor ligand family, and antibodies against the polypeptides, useful treating melanoma, sarcoma, and viral, bacterial, fungal infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (YUGG/) YU G.
(NIJJ/) NI J.
(ROSE/) ROSEN
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                                                        processing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 16; Fig 1; 29pp; English.
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                                       endokine alpha.
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  169
                                                           The present sequence represents the amino acid sequence of
AA;
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97US-00912227
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The invention relates to an isolated endokine alpha polypeptide (a member of the tumour necrosis factor family), appearing as ADC01858, the sequence of the endokine alpha polypeptide having an sequence encoded by the cDNA clone contained in ATCC Deposit number 97640 (A1) and the sequence of an epitope-bearing portion of the above polypeptides. Also included are the encoding nucleic acid (its homologues, complements or fragments where the fragment comprises at least 50 contiguous nucleotides), provided that the fragment is not from a region starting at nucleotide 26 and ending at nucleotide 476 of ADC01857), making a recombinant vector comprising the nucleic acid and an isolated antibody or antibody fragment that binds specifically to endokine alpha. The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chronic lymphocyte disorder; tumour; parasitic disease; autoimmune disease; lupus; arthritis; multiple sclerosis; inflammation; graft versus host disease; transplant rejection; skin allergy; bowel disease; wound; sepsis; Hodgkin's disease; chronic lymphocyte leukaemia; Burkitt's lymphoma; scleroderma; chronic active hepatitits; myasthenia gravis; psoriasis; autoimmune thyroiditis; Goodpasture's disease; asthma; Graves disease; cirrhosis; insulin dependent diabetes mellitus; Sjogren's syndrome; glomerulonephritis; hepatitis; Parkinson's disease; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated endokine alpha polypeptide AIDS, chronic lymphocyte disorder, common variable immunodeficiency, a tumor, parasitic disautoimmune disease, lupus, arthritis, multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; SEQ ID NO 2; 90pp; English.
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28-MAY-1999;
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99US-00345790.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00912227.
99US-0122099P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96US-0024058P.
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2. .169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "This protein is specifically claimed
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RRESULT 6
AAY15817
ID AAY15817
XX AAY15817
XX AAY1
XX AAY1
XX AAY1
XX AAY1
XX DAI
DT 28-J
DT 28-J
DT WA1
XX AAY1
DT DT 28-J
DT 27-M
XX WO99
PD 27-M
XX PF 18-N
PR 18-N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arthritis, idiopathic thrombocytopenic purpura, multiple sclerosis, arthritis, idiopathic thrombocytopenic purpura, multiple sclerosis, chronic inflammation, acute inflammation, acute allograft rejection, graft versus host disease, transplant rejection, foetal resorption, faecal peritonitis, skin allergies, bowel disease, a wound, sepsis, ALL, flodgkin's disease, non-Hodgkin's lymphoma, chronic lymphocyte leukaemia, plasmacytomas, multiple myeloma, Burkitt's lymphoma, EBV-transformed disease, chronic myelogenous leukaemia, chronic hypergammaglobulinaemeia, autoimmune haematcological disorders, polychondritis, scleroderma, Wegener granulomatosis, dermatomyosisitis, chronic active hapatitis, myasthenia gravis, psoriasis, Steven-Johnson syndrome, idiopathic sprue, autoimmune thyroiditis, idiopathic Addison's disease, vitiligo, gluten-sensitive entercents, autoimmune active hapatitis, chronic active hapatitis, myasthenia gravis, psoriasis, Steven-Johnson syndrome, idiopathic sprue, autoimmune controlleria, chronic active hapatitis, myasthenia gravis, psoriasis, Steven-Johnson syndrome, vitiligo, gluten-sensitive entercents, autoimmune active hapatitis, chronic active hapatitis, chronic active hapatitis, myasthenia gravis, psoriasis, chronic active hapatitis, autoimmune hapathic sprue, autoimmune hapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          enferopathy, autoimmune neutropenias, pemphigus vulgaris, Goodpasture's disease, bullous pemphigoid, discoid lupus, dense deposit disease, endocrine ophthalmopathy. IBD, asthma, Graves disease, Sarcoidosis, cirrhosis, juvenile diabetes, insulin dependent diabetes mellitus, uveitis, autoimmune gastritis, lymphopaenias, olyarteritis nodosa, Sjogren's syndrome, Bechet's disease, Hashimoto's disease, primary myxedema, polymyositis, mixed connective tissue disease, interstitial keratoconjunctivitis sicca, vernal keratoconjunctivitis, hammilitial
                                                                                                                                                                                                                                                                                                                                         DNA19355; tumour necrosis factor homologue; receptor GITR; immunogen; antibody; apoptosis; mammalian cancer cell; tumour necrosis factor (TNF)-alpha secretion; primary T- cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lung fibrosis, glomerulonephritis, hepatitis, autoimmune haemolytic anaemia, contact sensitivity disease, Parkinson's disease, primary lateral sclerosis, silicosis, sarcoidosis, idiopathic pulmonary fibrosis, idiopathic hyper-eosinophilic syndrome, endotoxic shock, atherosclerosis, histamine-mediated allergic reactions, IgE-mediated allergic reactions, rheumatoid arthritis, plastic anaemia and myelodysplastic syndrome. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and protein are useful for treating an individual having a disorder chosen from AIDS, chronic lymphocyte disorder, common variable immunodeficiency, a tumour, parasitic disease, autoimmune disease,
                                                                                                                            27-MAY-1999.
                                                                                                                                                                                      WO9925834-A1
                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                            proinflammatory response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acid sequence of human DNA19355 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY15817 standard; protein; 177 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 169
      18-NOV-1997;
                                                                18-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence represents endokine alpha.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NPOFIS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
      97US-0065635P
                                                                   98WO-US024621.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 675; DB 7;
Pred. No. 2.2e-69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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RESULT 7
AAYO6646
ID AAYO
XX AAYO
AC AAYO
XX 26-C
XX PRO3
XX PRO3
XX PRO3
XX Linfl
X
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Best Local S
Matches 126
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                                        Modified-site
                                                                                                                                                                                          Domain
                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                     Domain
                                                                                                                                                                                                                                                                        Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRO364 ligand; tumour necrosis factor receptor; human; apoptosis; inflammation; antiinflammatory; NF-KB activation; autoimmune dise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY06646;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY06646 standard; protein; 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mammalian cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ashkenazi AJ,
Godowski PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRO364 ligand.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fig
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Mark MR;
                                52. .17
/note=
129
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                                                                                                                                                  /note= "transmembrane domain"
                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                       /note= "cytoplasmic domain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86pp; English
                                                                                                                     .177
"N-glycosylation
                                                                     "extracellular domain"
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Pred. No. 2.4e-69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease;
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Best Local S
                 PRO364; human; glucocorticoid-induced tumor necrosis factor; receptor; hdGIR; ligand; hGIRL; PRO175; tumour necrosis factor receptor; TNFR; human umbilical vein endothelial cell; HUVEC; cardiac hypertrophy; myocardial infarction; PGF_2alpha; trauma; cancer; angiogenesis; age related macular degeneration; antibody, periodontal disease; vascular-related drug targeting; atherosclerosis; hypertension; inflammatory vasculitides; Reynaud's disease; aneurysm; arterial restenosis; thrombophlebitis; tumor angiogenesis; lung; liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a putative ligand for PRO364 (see also AAY06605), a novel member of the tumour necrosis factor receptor family. The sequence was deduced from an isolated cDNA clone (see AAX87726). Hydropathy analysis suggests a type II transmembrane homology. The mol.wt. is 20,308. Identity is shown to human Apo-2L (19.8%), Fas/Apolligand (19.0%), TNF-alpha (20.6%) and lymphotoxin-alpha (17.5%). PRO364 is useful for modulating apoptosis, NF-KB activation and proinflammatory or autoimmune responses in mammalian cells
                                                                                                                                                                              08-MAY-2001
                                                                                                                                                                                                        AAB47056;
            fibrosis;
                                                                                                                                                  PR0175
                                                                                                                                                                                                                                  AAB47056 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 177
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N-PSDB; AAX87726.
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            neuropathy;
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                                                                                                                                                                                                                                                                                                                                                        PFEVRLYKNKDMIQTLTNKSKIQNVGG
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                                                                                                                                                                                                                                  protein;
            rheumatoid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 675; DB 2;
Pred. No. 2.4e-69;
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            arthritis
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В
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                                                                                                                                                                                                                                                                                                                 antagonists, are useful for vascular-related drug targeting or as therapeutic targets for the treatment or prevention of atherosclerosis, hypertension, inflammatory vasculitides, Reynaud's disease, aneurysms, arterial restensis, thrombophlebitis, tumor angiogenesis, gut protection or regeneration and treatment of lung or liver fibrosis, periodontal diseases, attraction of bone-forming cells, central and peripheral nervous system disease and neuropathies and rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents PRO175 polypeptide, which is a human gluco-corticoid-induced tumor necrosis factor ligand (hGITRL). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; Fig 5; 76pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Composition for diagnosing and treating cardiovascular, endothelial angiogenic disorders, comprises a PRO364 or PRO175 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JUL-2000; 2000WO-US018867.
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                                                                                                                                                                                                          Local Similarity
  112
                                             61
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                                   PFEVRLYKNKDMI QTLTNKSKI QNVGGTYELHVGDTIDLI FNSEHQVLKNNTYWGIILLA
                                                                                                                                   ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                          ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
  PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
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                                                                                                                                                                                  Score 675; DB 4;
Pred. No. 2.4e-69;
); Mismatches 0;
                                                                                                                                                                                                                                Length 177;
                                                                                                                                                                                       Indels
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Gurney AL, Hillan KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiarthritic; antiinflammatory; antianaemic; immunosuppressive;
antithyroid; antidiabetic; neuroprotective; hepatotropic; viruci
dermatological; antipsoriatic; antiasthmatic; antiallergic;
                    The present sequence is that of PRO175 (UNQ149), a novel human immunomodulator protein (20 kDa, pl 8.08) showing homology to tumour necrosis factor family members. The invention provides polynucleotides (see AAF30050-62) encoding novel human PRO proteins (see AAB20108-20) including PRO175. Claimed compositions comprising these proteins or their agonists are useful for increasing infiltration of inflammatory cells into a tissue of a mammal, stimulating or enhancing an immune response in a mammal, or increasing the proliferation of T-lymphocytes in a mammal in response to an antigen. Claimed compositions comprising the PRO polypeptide or its antagonist have the opposite effect. A claimed method for treating an immune related disorder, such as a T cell disorder, involves administering the PRO polypeptide, an agonist antibody or an antagonist antibody. The disorder is selected from systemic lupus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New PRO polypeptides, nucleic acids and (ant)agonists, diagnosing and treating immune-related disorders, such sclerosis, rheumatoid arthritis and diabetes.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 20; Fig 4; 127pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-1999;
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Pitti RM,
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chyroiditts, diabetes mellitus, immune-mediated renal disease,

chyroiditts, primary biliary cirrhosis, granulomatous hepatitis,

chyroiditts, primary biliary compathy, Whipple's disease,

chyroiditts, disease, gluten-sensitive enteropathy, Whipple's disease,

chyroiditts, atopic dermatitis, allergic diseases (such as asthma,

chyroiditts, atopic dermatitis, food hypersensitivity and

chyroiditts, atopic dermatitis, food hypersensitivity and

chyroiditts, atopic diseases of the lung and transplantation

chyroiditts, atopic diseases of the lung and transplantation

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Best Local :
         23-JUN-1999;
20-JUL-1999;
26-JUL-1999;
01-SEP-1999;
                                                                                                                                                                                             12-MAR-1999;
14-MAY-1999;
02-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; angiogenesis-associated protein; PRO; endothelial cell growth cardiack hypertrophy; cardiovascular disorder; endothelial disorder; angiogenic disorder; atherosclerosis; osteoporosis; hypertension; myocardial infarction; diabetic retinopathy; rheumatoid arthritis; Crohn; disease; psoriasis; endometriosis; ulcer; wound healing; carricular dispersions of the control of the c
                                                                                                                                                                                                                                                                                                                                              08-MAR-1999;
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99US-0144758P
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Pred. No. 2.4
0; Mismatches
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No. 2.4e-69;
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The invention relates to novel human angiogenesis-associated proteins designated PRO proteins (AAB53064-B53097), and to nucleic acids encoding PRO proteins. The invention also relates to vectors and host cells comprising a PRO nucleic acid, the recombinant production of a PRO protein, PRO antibodies specific for a PRO protein, fusion protein, and compounds which inhibit the expression of a PRO gene. The invention additionally encompasses methods of identifying modulators of pRO expression or activity; diagnosing a cardiovascular, endothelial or angiogenic disorder, or a susceptibility to such a disorder by detecting mutations in a PRO gene, or the expression level of a PRO gene within a particular tissue; treating a cardiovascular, endothelial or angiogenic disorder to the expression level of a PRO gene within a particular tissue; treating a cardiovascular, endothelial or angiogenic disorder is the administration of a DRO protein and particular tissue; treating a cardiovascular, endothelial or angiogenic disorder is the administration of a DRO protein and the protein administration of a DRO protein and administration and administration of a DRO protein and administration and administration of a DRO protein and a DRO protein and a DRO protein and administration and 
                                                                                                                                                                                                                                                      PRO agonist or antagonist; a retroviral gene therapy vector comprising a PRO nucleic acid; and methods of inhibiting or stimulating endothelial cell growth, cardiac hypertrophy or PRO-induced angiogenesis via the administration of a PRO protein, or an agonist or antagonist thereof. PRO nucleic acids, PRO proteins, antibodies against PRO proteins, PRO agonists and PRO antagonists may be used as therapeutic agents to treat cardiovascular, endothelial or angiogenic disorders, such as atherosclerosis, osteoporosis, myocardial infarction, hypertension, diabetic retinopathy, rheumatoid arthritis, Crohn's disease, psoriasis, endometriosis, ulcers, wounds, cancer, Alzheimer's disease, Huntington's
                                                 endometriosis, ulcers, wounds, cancer, Alzheimer's disease, Huntington's disease, or stroke. PRO nucleic acids are additionally useful in the recombinant production of PRO proteins, as hybridisation probes to screen libraries to isolate cDNAs with sequence identity to PRO proteins, to map genes encoding PRO proteins, to analyse genetic disorders, and in gene therapy. PRO nucleic acids can also be used to produce transgenic animals useful for the development and screening of potential therapeutic agents. The present sequence represents a PRO protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-SEP-1999;
15-SEP-1999;
15-SEP-1999;
05-OCT-1999;
30-NOV-1999;
02-DEC-1999;
02-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 particular tissue; treating a cardiovascular, endothelial or angiogenic disorder via the administration of a PRO protein, PRO nucleic acid, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid for producing a PRO polypeptide, analyzing genetic disorders and treating cardiovascular, endothelial or angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disorders,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Paoni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ashkenazi AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    such as atherosclerosis, wounds or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pitti RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US020594.
99WO-US021090.
99WO-US021547.
99WO-US023089.
99WO-US028313.
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99WO-US028564.
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Hillan KJ,
tanabe CK, W
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Kuo SS, Mark MR, Mar
Mood WI;
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ark MR, Marsters SA;
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Sequence 177
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Best Local S
Matches 126
                                                                                                                  Local Similarity
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   121
                                                                                                       126;
                                           61
                                                                52
                                                                                    μ
                                                                           ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLBILQNGLYLIYGQVAPNANYNDVA
   NPQFIS 126
                                PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                            ETAKEPCMAKFGPLPSKWOMASSEPPCVNKVSDWKLEILONGLYLIYGQVAPNANYNDVA
                                                                                                        Conservative
                                                                                                                 100.0%;
                                                                                                       <u>.</u>
                                                                                                               Score 675; DB 4;
Pred. No. 2.4e-69;
                                                                                                       Mismatches
                                                                                                                        Length
                                                                                                      Indels
                                                                                                                        177;
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                                                                                                     Gaps
                                         120
                                                             111
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RESULT 11
AAB47287
ID AAB47
XX AAB47
XX AAB47
XX AAB47
XX AAB47
XX PRO;
XX PRO
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30-NOV-1999;
05-JAN-2000;
24-FEB-2000;
15-MAR-2000;
30-MAR-2000;
17-MAY-2000;
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The sequences given in AAB47287-90 show PRO polypeptides. PRO175 shows type II transmembrane protein typology, and portions of PRO364 show homology to members of the tumour necrosis factor (TMP) family, thereby indicating that it may be a novel member of the TMP family. PRO175 and PRO185 stimulated heart hypertrophy. PRO cDNA's may be used to identify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                    Isolated PRO polypeptide useful in treating and diacardiovascular, endothelial or angiogenic disorder mellitus, myocardial infarction, arthritis.
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                                                                                                                                                   Claim 56; Fig
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                                                                                                                                                                                                                                                                                                                                                          i AJ, L.
^T. Mark MR,
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2000WO-US005004.
2000WO-US006884.
2000WO-US008439.
2000WO-US013705.
2000WO-US014941.
2000WO-US014941.
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                                                                                                                                                   2; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         Baker
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161. .16
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/label= N-myristoylation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     129.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /label= N-linked glycosylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'label= Transmembrane domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                          INC.
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                                                                                                                                                                                                                                                                                                                                                                                    Marsters
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                                                                                                                                                                                                                                                                                                                                                                                 Ferrara N, Godowski PJ, (
sters SA, Paoni NF, Pitti
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                                                                                                                                                                                                                                                             diagnosing
                                                                                                                                                                                                                                 e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                       Gurney AL;
i RM, Wood
                                                                                                                                                                                                                                                             a
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                                                                                                                                                                                                                                      diabetes
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RESULT 12
AABSULT 12
AABSU XX
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XX 21-M2
XX 21-M2
XX 1-M2
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XX Humar
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Matches
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02-DEC-1999;
16-DEC-1999;
05-JAN-2000;
06-JAN-2000;
11-FEB-2000;
18-FEB-2000;
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24-FEB-2000;
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20-JUL-1999;
26-JUL-1999;
28-JUL-1999;
01-SEP-1999;
30-NOV-1999;
30-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; PRO; cardiant; antiangiogenic; antiarteriosclerotic; hypotensive vasotropic; antirheumatic; antiarthritic; antiinflammatory; cytostatic; vulnerary; antianginal; gene therapy; cardiovascular disease; endothelial disorder; angiogenic disorder; cancer; periodontal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amylotrophic lateral sclerosis, and stroke
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 675; DB 4; ilarity 100.0%; Pred. No. 2.4e-69; Conservative 0; Mismatches 0:
  99WO-US030095.
2000WO-US000376.
2000WO-US000376.
2000WO-US003365.
2000WO-US004341.
2000WO-US004342.
2000WO-US004342.
2000WO-US004343.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-US013705
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                                                                                                                                                                                                                                              99US-0146222P.
99WO-US020111.
99WO-US028313.
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99US-0141037P.
99US-0144758P.
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                                                                                                                                                                                             99WO-US028565.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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AAU81953
ID AAU81
XX AAU81
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XX Humar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Simi
Matches 126;
   Human; PRO; antiinflammatory; ophthalmological; vasotropic; retinal cell injury; ocular disease; retinitis pigmentosa; macular degeneration; retinal detachment; retinal tear; retinopathy; retinal degenerative disease; macular hole; degenerative myopia; acute retinal necrosis syndrome; traumatic chorioretinopathy; Purtscher's retinopathy; oedema; ischaemic condition; retinal vision occlusion; collagen vascular disease; thrombocytopaenic purpura; uveitis; retinal vasculitis; Eales disease systemic lupus erythematosus; environmental trauma.
                                                                                                                                                                            Human PRO175
                                                                                                                                                                                                           09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is one of seventeen novel PRO polypeptides. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating cardiovascular, endothelial or anglogenic disorders in a mammal. Examples of these disorders include cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restencesis, Reynaud's disease, rheumatoid arthritis, angina, myocardial infarctions, thrombophlebitis and lymphangitis. The PRO polypeptides and antagonists are also used to prevent tumnour angiogenesis and for treating periodontal diseases. They are also used to stimulate wound healing and antibodies are useful for diagnosing a cardiovascular, endothelial or angiogenic disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-MAR-2000;
10-MAR-2000;
15-MAR-2000;
21-MAR-2000;
30-MAR-2000;
                                                                                                                                                                                                                                                                        AAU81953 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 71; Fig 2; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ashkenazi
Goddard A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disorders in a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Paoni NF, Pitti RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GETH )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   iagnosis
                                                                                                                                                                                                                                                                                                                                                         172
                                                                                                                                                                                                                                                                                                                                                                                      121
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                                                                                                                                                                                                                                                                                                                                                                                       NPQFIS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                   PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A Godowski
                                                                                                                                                                                                                                                                                                                                                         NPQFIS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ETAKBPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acids and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000WO-US005841.
2000WO-US006319.
2000WO-US006884.
2000WO-US007532.
2000WO-US008439.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA;
                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            r KP,
ki PJ, Gurne,
יש. Watanabe CK,
                                                                                                                                                                                                                                                                      protein; 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         encoding PRO polypeptides which are useful in of cardiovascular, endothelial or angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 675; DB 4;
Pred. No. 2.4e-69;
                                                                                                                                                                                                                                                                        B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , Gerber H, Gerritsen
, Kuo SS, Mark MR, Ma
Williams PM, Wood WI;
                         vasculitis; Eales disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Marsters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ME.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>,</u>
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ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEIIQNGLYLIYGQVAPNANYNDVA ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA

Query Match Best Local S Matches 126

Similarity

100.0%; ilarity 100.0%; Conservative

0,

Score 675; DB 5; Pred. No. 2.4e-69; ; Mismatches 0;

Length 177; Indels

0,

Gaps

111 60 Sequence

177

A

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CC The invention relates to promoting the survival of retinal cells, or C delaying or preventing retinal cell injury or death, by contacting the CC retinal cells with the polypeptide such as PRO175, PRO220, PRO216, PRO216, PRO316,                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JAN-2000; 2
06-JAN-2000; 2
11-FEB-2000; 2
18-FEB-2000; 2
22-FEB-2000; 2
24-FEB-2000; 2
24-FEB-2000; 2
15-MAR-2000; 2
15-MAR-2000; 2
30-MAR-2000; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JUL-1999;
13-SEP-1999;
15-SEP-1999;
29-NOV-1999;
30-NOV-1999;
01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Promoting survival of retinal cells, or delaying or preventing retinal cell injury or death, by contacting retinal cells with PRO175, 220, 216, 243, 306, 346, 322, 536, 943, 840, 828, 826, 1068 or PRO1132 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kljavin
Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUL-2000;
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)B; ABK28562.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44; Fig 2; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENENTECH
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99WO-US028313

99WO-US028313

2000WO-US000219

2000WO-US000376

2000WO-US003565

2000WO-US004414

2000WO-US005044

2000WO-US005044

2000WO-US0050441

2000WO-US0050441

2000WO-US0050441

2000WO-US00505
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US021090
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mark MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Goddard A, Godowski PJ, Gurney
Mark MR, Marsters SA, Pitti RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gurney AL;
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ARBSULT 14
ABR42323
ID ABR42
XX ABR422
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XX III-AU
XX III-AU
XX Humar
XX Humar
XX Humar
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XX Hom
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                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is the protein sequence of human GITRL polypeptide. The invention relates to compositions comprising heterotrimeric complexe of tumour necrosis factor (TNF) ligand family members, and their use in the detection, prevention and treatment of disease. In preferred embodiments, the heterotrimeric complex comprises full-length or extracellular portions of GITRL together with full-length or extracellular portions of other TNF ligand family members. The heterotrimeric complexes of the invention are useful for treating an heterotrimeric complexes of the invention are useful for treating an entirement defense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New heteromultimeric complex having a first polypeptide member of the tumor necrosis factor (TNF) ligand family, and a second different mem of TNF ligand family, useful for treating cancer, osteoporosis or an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hilbert DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUL-2001; 2001US-0307838P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-MAY-2003
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                                                                                                                                                                                                                                                                                                                                               Sequence 177
                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune disease, cancer or osteoporosis, and particularly for inhibiting cancer cell proliferation, increasing B cell proliferation, inducing apoptosis of T cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     112
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DB; ACC57909.
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                                                                                                                                                                                                                                    126;
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                                                     PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                    ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 382-383; 388pp; English.
                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen CA;
                                                                                                                                                                                                                                                                                                                                                     AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour necrosis factor; ligand; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteopathic.
                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                            Score 675; DB 6;
Pred. No. 2.4e-69;
                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                          Length 177;
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                                                                                                                                                                                                                                       Gaps
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RESULT 15
RABF60544
IID ABP60
XX ABP60
XX ABP60
XX APR10
DT 28-MA
XX APR11
KW APR11
KW APR11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human, disease or disorder such as autoimmune disease, and graft versus host disease (GVHD). The autoimmune disease is systemic lupus erythematosus, rheumatoid arthritis or Sjogren's syndrome. The antibody is useful for detecting, diagnosing, prognosing, treating, preventing on ameliorating a disease or disorder associated with aberrant APRIL or APRIL receptor expression or aberrant function of APRIL or APRIL receptor. The disease or disorders includes autoimmune and inflammatory receptor. The disease or disorders includes autoimmune and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antiinflammatory, antirheumic, antiarthritic, cytostatic, antianaemic, antiallergic, antiasthmatic, neuroprotective, ophthalmological, tuberculostatic, antidiabetic, antidiabetic, antidiateric, antidiabetic, antidiateric, and haemostatic activity antiarteriosclerotic, vascotropic, thyromimetic, and haemostatic activity. The antibody or its fragment are useful for treating, preventing or manipulating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as Non-Hodgkin's lymphoma or multiple myeloma in ameliorating cancer such as 
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                                                                                                                                              gravis, Hashimoto's disease, immunodeficiency syndrome, Bruton's disease, infectious diseases (e.g. acquired immunodeficiency syndrome (AIDS)), and proliferative disorders (e.g. leukemia). The present sequence represents
                                                                                                                                                                                                                                                                                                                                                            disorders such as autoimmune neutropenia, haemolytic anaemia, dermatitis, asthma, allergic encephalomyelitis, myocarditis, multiple sclerosis, uveitis, tuberculosis, diabetes mellitus, psoriasis, cancer of the immune
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Search completed: November 2, 2004, 12:55:27 Job time : 157 secs	172 NPQFIS 177	121 NPQFIS 126	112 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSBHQVLKNNTYMGIILLA 171	61 PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120	52 ETAKEPCMAKFGPLÞSKWQMASSEÞPCVNKVSDWKLEILQNGLYLIYGQVAÞNANYNDVA 111	1 ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLBILQNGLYLIYGQVAPNANYNDVA 60	~	Query Match 100.0%; Score 675; DB 6; Length 177; Best Local Similarity 100.0%; Prod No. 2 40-60.

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Sequence 2, Appli
Sequence 6, Appli
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Sequence 7, Appli
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Sequence 7, Appli
Sequence 7, Appli
Sequence 70, Appli
Sequence 71, Appli
Sequence 71, Appli
Sequence 77, Appli
Sequence 77, Appli
Sequence 78, Appli
Sequence 73, Appli
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1 ETAKEPCMAKFGPLFSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA

Query Match 100.0%; Score 675; Best Local Similarity 100.0%; Pred. No. 2. Matches 126; Conservative 0; Mismatches	US-08-912-27-2 US-08-912-27-2 Sequence 2, Application US/08912227 Patent No. 5998171 GENERAL INFORMATION: APPLICANT: Yu, Guo-Liang APPLICANT: Ni, Jian APPLICANT: Rosen, Craig A. TITLE OF INVENTION: Human Endokine Alpha CORRESPONDENCE ADDRESS: ADDRESSE: STERNE, KESSLER, GOLDSTEIN & STREET: 1100 NEW YORK AVE., NW, SUITE 6 COUNTRY: USA ZIF: 20005-3934 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PAPLICATION DATA: APPLICATION NUMBER: US/08/912,227 FILING DATE: Herewith CLASSLETICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: US/08/912,227 FILING DATE: 16-AUG-1996 ATTORNEY/AGENT INFORMATION: NAME: Steffe, Eric K. REGISTRATION LOTA: APPLICATION INFORMATION: TELEPHONE: 202-371-2600 TELEPHONE: 202-371-2600 TELEPHONE: 202-371-2540 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 169 amino acids TYPE: amino acids TYPE: amino acids TYPE: protein US-08-912-227-2	.5 10.3 984 2 US-0 .5 10.3 984 4 US-0 .5 10.3 984 4 US-0 .5 10.3 989 4 US-0 .5 10.3 989 3 US-0 .5 10.3 1013 4 US-0 .5 10.1 1604 3 US-0 .5 10.1 1604 3 US-0 .5 10.1 3443 2 US-0 .5 10.0 211 4 US-0 .5 10.0 211 4 US-0 .5 10.0 289 4 US-0 .5 10.0 289 4 US-0 .5 10.0 289 4 US-0	8 69.5 10.3 984 2 US-08
2.DB 2; Length 169; 2.2e-75; 2.2e-75; Indels 0; Gaps 0;	ion #1.30	Sequence 12 Sequence 12 Sequence 12 Sequence 12 Sequence 12 Sequence 2, Sequence 18 Sequence 18 Sequence 19 Sequence 2, Sequence 16 Sequence 16 Sequence 16 Sequence 70 Sequence 70 Sequence 70 Sequence 70 Sequence 65 Sequence 38 Sequence 38	Sequence 12

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                                                                       ; MOLECULE TYPE: US-09-513-584-2
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Patent No. 6406867
GENERAL INFORMATION:
Query Match 100.0%; Score 675; DB 4;
Best Local Similarity 100.0%; Pred. No. 2.2e-75;
Matches 126; Conservative 0; Mismatches 0;
                                                                                                                                                         TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/09/513,584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 6:
FILING DATE: 16-AUG-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                         REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 26-FEB-
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Human Endokine Alpha and Methods TITLE OF INVENTION: of Use NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                             FILING DATE: 01-JUL-1999
ATTORNEY/AGENT INFORMATION:
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APPLICANT: Rosen, Craig A.
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ZIP: 20005-3934
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FO
ADDRESSEE: P.L.L.C
STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                      TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/912,227 FILING DATE: 15-AUG-1997
                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                 REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 28-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                          LENGTH:
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                                                                                                                                            169 amino acids
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Patent No.
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Matches 126; Conserv
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION IMFORMATION:
TELEPHONE: 202-371-2600
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                        TYPE: ami TOPOLOGY:
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5. 6521742
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)GY: linear
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                                                                                                PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
                                          NPOFIS 126
                                                                            PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                                                                                                   ETAKEPCMAKFGPLPSKWOMASSEPPCVNKVSDWKLEILONGLYLIYGOVAPNANYNDVA
         NPQFIS 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DC
                                                                                                                                                                                                                                                                                                                                                             169 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yu, Guo-Liang
Ni, Jian
                                                                                                                                                                                                                                                                                                                                                                                                              202-371-2540
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                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                        protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Endokine Alpha
                                                                                                                                                                                                                    100.0%; Score 675; DB 4;
100.0%; Pred. No. 2.2e-75;
tive 0; Mismatches 0;
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NW, SUITE 600
                                                                                                                                                                                                                                                      Length 169;
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Gaps

163

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US-09-248-796A-20671

Sequence 20671, Application US/09248796A

Patent No. 6747137

; GENERAL INFORMATION:
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                                                                                                                                                     RESULT 5
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Sequence 3959, Application US/09107532A

Patent NO. 5833275

GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-107-532A-3959
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 19; Conserv
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (781)893-5007
TELEPAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 3959:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...1221
SEQUENCE DESCRIPTION: SEQ ID NO: 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/051571
FILING DATE: JULY 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: ARINIELLO, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1398
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL: YES ORIGINAL SOURCE:
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                 575
                                                                                                                                                                                                                                                                                             515
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                                                                                                                                                                                                                                                   106 QVLKNN 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                           51 APNANYNDVAPFEVRLYKNKDMIQTLTNKS-----KIQNVGGTYELHVGDTIDLIFNSEH 105
                                                                                                                                                                                                                                                                                     APNGYQLDSSEFSITVKENQNVTTTRINKKSTGSVEIEKIGDLDGLGLPNVVFTIYNSDN 574
                                                                                                                                                                                                               KVVKDN 580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 1221 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: <Unknown>
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COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                         12.5%; Score 84.5; DB 28.8%; Pred. No. 0.45;
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                                                                                                                                                                                                                                                                                                                                                                          19; Mismatches
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AND AMINO ACID SEQUENCES RELATING TO
PAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                            4;
                                                                                                                                                                                                                                                                                                                                                                        23;
                                                                                                                                                                                                                                                                                                                                                                                                               Length 1221;
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Query Match
Best Local Similarity
Warches 31; Conserva
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PRIOR FILING DATE: 1998-02-13
PRIOR PRIOR PAPELICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 26671
LENGTH: 217
ORGANISM: Candida albicans
FEATURE:
                                                                                      US-08-496-944-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-496-944-2
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APPLICANT: Law, Marcus I
APPLICANT: Dietz, Jon M
                                                                                                                                                                        REFERENCE/DOCKET NUMBER: INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/248,796A CURRENT FILING DATE: 1999-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 107196.132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: UNSURE LOCATION: (15)
OTHER INFORMATION:
                                                                                                 TOPOLOGY: li
                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Elmer, James Scott
REGISTRATION NUMBER: 36,129
                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF ANDRESS: 8
CORRESPONDENCE ADDRESS:
ANDRESSEE: CIBA-Geigy
                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: / STREET: / Hawthorne
                                                                                                                                         TYPE:
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: CIBA-----ADDRESSEE: 7 Skyline Drive
                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10532
                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 PSKWQMASS--EPP-----CVNKVSDWKLEILQNGLYLIYGQVAP----NANYNDVAPF
                                                                                                                                       amino acid
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                                                                                                                                                        2763 amino acids
                11.3%;
ilarity 25.8%;
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                                                                                                     protein
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                                                                                                                                                                                                                                                                                   435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11.4%;
27.2%;
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                Score 76; DB:
Pred. No. 17;
18; Mismatches
                                                                                                                                                                                                          CGC 1814
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Pred. No. 0.3;
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                                                                                                                                                                                                                                                                                                                                                 Version
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                                                Length 2763;
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18;

Indels

20;

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PCT-US93-02475-6
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                                                                                                             Matches
                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/.
APPLICATION NUMBER: US 07/.
PILING DATE: 12 March 1992
ATTORNEY/AGENT INFORMATION:
NAME: Donald G. Lewis
REGISTRATION NUMBER: 28636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (619) 554-6312 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wisnieski, Bernadine J. TITLE OF INVENTION: Tumor Necrosis TITLE OF INVENTION: Ion Channel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: BJ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 157 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                      PUBLICATION INFORMATION:
AUTHORS: McGraw, R. A
                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM:
SOFTWARE: Word Pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 199304
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STATE: California
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                                                                                                                                                                                          PAGES: 5564
DATE: 1990
RELEVANT RESIDUES IN SEQ ID NO:
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                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                  VOLUME:
                                                                                                                                                                                                                                                                     JOURNAL:
                                                                                                                                                                                                                                                                                                                        AUTHORS:
                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: Tumor Necrosis Factor (feline)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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     81
                                        42
                                                                       30 KVSDWKLEILQNGLYLIYGQV-----APNANY---NDVAPFEVRLYKNKDMIQTLINKS 80
                                                                                                                                                                                                                                                                                                                                                                                                                                           AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VSRWYKNRKESLKTDSLESFRNK-----
                                                                                                                                                                                                                                                                                   S: McGraw, R. A., Coffee, B.W., Otto, S: C.M., Drews, R.T. and Rawling, C.A. Gene Sequence of Feline Tumor Necrosis Factor '.
     KIQN--
                                      BLTDNQLKVPSDGLYLIYSQVLFTGQGCPSTHVLLTHAISRFAVSYQTKVNLLSAI--KS 99
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                                                                                                             Conservative
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                                                                                                           11.0%; Score 74; DB 24.4%; Pred. No. 0.4! tive 21; Mismatches
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         -VGGTYELHVGDTID-
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                                                                                                                                              DB 5; Length 157;
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                                                                                                             32;
                                                                                                               Indels
         -LIFNSEHQVLKNNTYWGI 116
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                                                                                                               Gaps
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APPLICANT: Reed, John C.
APPLICANT: Sato, Takaaki
TITLE OF INVENTION: CD40
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/349,357
PILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1203
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 235
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                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Diego
STATE: California
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                                                                                                                                                                       CLASSIFICATION: 530
                                                                                                                                                                                                               APPLICATION NUMBER:
  TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         33; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    74
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                                                                                                                                                                                                                                                                                                                                                                                                                         E: Campbell and Flores
4370 La Jolla Village I
                                                                                                                                                                                                                                                                                                                                                                     USA
(619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31; Indels
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US-08-349-357-2
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                                                                                                                                                            US-08-349-357-2
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Matches
                                                                                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Reed, John C.
APPLICANT: Sato, Takaaki
TITLE OF INVENTION: CD40 Associated Proteins
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: C
                                                                                                                                                                                       TOPOLOGY:
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                                                                                                                                                                                                               LENGTH: 543 amino acids
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 282
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                                                                            21 ASSEPPCYNKVSDW-----KLEILQNGLYLIYGQVAPNAN----YNDVAPFEVRLYKNK 70
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                      71 DMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLK 109
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EMLR--NNESKI
                                                 ASSAVOHVNLLKEWSNSLEKKVSLLON-----ESVEKNKSIQSLHNOICSFEIEIEROK 281
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                                                                                                                  11.0%;
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                                                                                                       17; Mismatches
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                                                                                                                  Score 74; DB 3; Pred. No. 2.7;
LHLQRVID-
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                                                                                                     24; Indels
                                                                                                                              Length 543
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CURRENT FILING DATE: US/09/645,926A
CURRENT FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: 60/151,250
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7
LENGTH: 567
TYPE: PRT
ORGANISM: Homo sapiens
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US-09-645-926A-7
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PRIOR FILLING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILLING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILLING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILLING DATE: 1998-08-13
PRIOR FILLING DATE: 1998-08-13
PRIOR FILLING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 15588
LENGTH: 289
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Patent No. 6482411
GENERAL INFORMATION:
APPLICANT: AHUJA, SEEMA
APPLICANT: BONEWALD, LYNDA
TITLE OF INVENTION: CD40 LIGAND AND CD40 AGONIST COMPOSITIONS AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y Match 11.0%; Score 74; DB 4; Length 567;
Local Similarity 28.3%; Pred. No. 2.9;
hes 28; Conservative 17; Mismatches 24; Indels
114 WGIILLANPQFI 125
                                                   197 RKSIHTLTNNYKKFVONIKTEADKYHWLTIIRANTTKIYQYIYSKLFYNSE-
                                                                                                                                                       143 IAINKPSSISLATDLKYQVIPYFNGIY---
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                                                                                                      70 KDMIQTLTNKSK--IQNV--------GGTYELHVGDTIDLIFNSEHQVLKNNTY 113
                                                                                                                                                                                                       20 MASSEPPCVNKVSDWKLEILQ--NGLYLIYGQVAPNANYNDVAPFEVRLY------KN
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                                                                                                                                                                                                                                                     10.8%; Score 73; DB 4; ilarity 22.7%; Pred. No. 1.4; Conservative 23; Mismatches 3
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Patent No. 6538176
GENERAL INFORMATION:
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LENGTH: 609
TYPE: PRT
ORGANISM: Xenopus laevis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application PC/TUS9302475 GENERAL INFORMATION:
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Best Local (
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CURRENT FILLING DATE: 1999-09-15
CURRENT FILLING DATE: 1999-09-15
NUMBER OF SEQ ID NOS: 22
SOFTWARE: FASTSEQ for Windows Version 3.0
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TITLE OF INVENTION: Maize Replication Protein A and
FILE REFERENCE: 5718-59
                TELEFAX: (619) 554-63
INFORMATION FOR SEQ ID NO:
                                                                     REFERENCE/DOCKET NUMBER: BUTTELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3
COMPUTER: VE System 386
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                          SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                               APPLICATION NUMBER: US 07/852,625
FILING DATE: 12 March 1992
TTORNEY/AGENT INFORMATION:
NAME: Donald G. Lewis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 10.7%; Local Similarity 24.3%;
                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                REGISTRATION NUMBER:
                                                                                                                                                                                                                                                     APPLICATION NUMBER: POFILING DATE: 19930412
                                                       TELEPHONE:
                                                                                                                                                                                                                                 CLASSIFICATION:
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WENTION: Tumor Necrosis Factor with Modified
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                                                         (619)
                                  554-6312
                                                         554-2421
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Pred. No. 5.
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US-09-710-279-606
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                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Best Local Similarity
Matches 33; Conserv
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                                                                                                              Matches
                                                                                                                              Best Local Similarity
                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                                    FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: amino acid sequence
                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                LENGTH: 620
TYPE: PRT
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AUTHORS: Pauli, U. Beutler, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: Tumor Necrosis Factor (porcine)
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DATE: 1989
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                                    450 NEWORI-----MMSQQRFKVLANGFVELYGYVTINVDR-----DVTVGINENIVON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87 YOTKVNLLSAI--KSPCORETPEGAEAKPWYEPIYLGGVFQLEKDDRLSAEINLPDYLDF 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27 QWQSGYANALLANGVKLKDNQLVVPTDGLYLLYSQVLFRGQGCPSTNVFLTHTISRIAVS 86
                                                                       16 SKWOMASSEPPCVNKVSDWKLEILONGLYLIYGOVAPNANYNDVAPFEVRLYKNKDMIQT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   66 LYKNKDMIQTLTNKSKIQN---
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76 L----TNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQ 123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Porcine Tumor Necrosis Factor-': Cloning with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                185-191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09710279
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                                                                                                           10.5%; Score 71; DB 4; Length 620 23.0%; Pred. No. 7.8; tive 20; Mismatches 33; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A blank residue designated by wide is inserted after residue No. 7 of porcine TNF and the sequence numbering is augmented by 1 starting with residue No. 8 in order to maximize tarting with human TNF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19; Mismatches
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Pred, No.
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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/cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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    5795
543
543
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177
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113
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_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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US-10-080-455-1
US-10-16-378-16
US-10-151-882-39
US-10-218-547-40
US-10-310-793-44
US-10-338-083-17
US-10-611-399-17
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                   US-09-815-242-12610
US-09-757-041-2
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Sequence 2, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 16, Appl
Sequence 39, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 47, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 12610, A
Sequence 2, Appli
Sequence 35, Appli
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10.4	10.4	٠	10.5	•	10.5	10.5	10.6	10.6	10.6	10.6	10.6	10.7	10.7	10.7	10.7	10.7	10.7	10.8	10.8	10.8	٠	10.8			٠	11.0	11.0	11.0	11.0	11.0	11.0
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US-10-297-942-2	US-10-437-963-159310	-10-474-	US-10-282-122A-74118	US-09-815-242-13477	US-10-282-122A-71355	-10-295-	-10-295-027-	US-10-154-759-17	-10-154-759-	-10-154-759-	-10-154-759-	-10-437-	-10-372-	0-375-553	-10-371-558-	-10-372-686-	-10 - 424	-10-295-074-	-10-295-074-5	-10-295-074-	-10-	0-424-599-23192	-10-437-963-12546	US-10-437-963-184313	0-042-865-1	0-004-378A-3	US-10-116-275-173	US-08-813-323A-2	-10-207-65	S-10-242-	US-08-813-323A-1
Sequence 2, Appli	15931		Sequence 74118, A	Н	e 7135	42, 1	280, Ap	17, App	e 15,		16,	1535	9	6, Appl	6, Appl	6, 2	209463	53,	59,		49,	231929	equence 125464	184	166		e 173	2, Appli	e 103, Ap	e 7,	Sequence 1, Appli

ALIGNMENTS

US-09-345-790-2

Sequence 2, Application US/09345790 Patent No. US20020099198A1 GENERAL INFORMATION:

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CLASSIFICATION:
PRIOR APPLICATION DATA:
PRICATION NUMBER: 08/912
PILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,66
                                NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 148
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/345,790
                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER,
STERET: 1100 NEW YORK AVE.,
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Human Endokine Alpha NUMBER OF SEQUENCES: 10
                     TELEPHONE:
                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 20005-3934
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 126; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10136511 Publication No. US20020168729A1
                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/122,099
PRIOR FILING DATE: 1999-02-26
PRIOR APPLICATION NUMBER: 08/912,227
PRIOR FILING DATE: 1997-08-15
PRIOR APPLICATION NUMBER: 60/024,058
PRIOR FILING DATE: 1996-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 09/513,584
PRIOR FILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: 09/345,790
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 60/136,788
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1999-05-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE OF INVENTION: Human Endokine Alpha and Methods of Use FILE REFERENCE: 1488.0470007/EKS/PSC CURRENT APPLICATION NUMBER: US/10/136,511 CURRENT FILING DATE: 2002-05-02
                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 169
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hes 126; Conserva
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121 NPQFIS 126
                                           104 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
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                                                                                    PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
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Pred. No. 3.1e-67;
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Publication No. US20020146389A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Godowski, Paul J.
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LENGTH: 177
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Publication No.
                                                                    SEQ ID NO 16
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                                                                                        CURRENT APPLICATION NUMBER: US/10/116,378

CURRENT FILING DATE: 2002-04-04

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/247,225

PRIOR FILING DATE: EARLIER FILING DATE: 1999-02-09

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/074,087

PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-09

NUMBER OF SEQ ID NOS: 31
                                                                                                                                                                                                                                                              APPLICANT: Goddard, Audrey
APPLICANT: Gurney, Austin
APPLICANT: Marsters, Scot A.
APPLICANT: Pitti, Robert M.
APPLICANT: Wood, William
TITLE OF INVENTION: NOVEL TUMOR NECROSIS FACTOR RECEPTOR HOMOLOG
TITLE OF INVENTION: NCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P1206R1
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ashkenazi, Avi J. APPLICANT: Goddard, Audrey APPLICANT: Gurney, Austin
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CURRENT EPPLICATION NUMBER: US/10/080,455
CURRENT FILING DATE: 2002-02-22
PRIOR APPLICATION NUMBER: 09/195,368
PRIOR FILING DATE: 1998-11-18
PRIOR PILING DATE: 1998-11-18
PRIOR APPLICATION NUMBER: US 60/069,661
PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: US 60/065,635
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APPLICANT:
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TYPE: PRT
ORGANISM: Homo sapiens
ORGANISM: Homo sapiens
                      LENGTH: 177
TYPE: PRT
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Pitti, Robert M.
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RESULT 6
US-10-218-547-40
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US-10-151-882-39
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 39
LENGTH: 177
TYPEF.
                                                                                                                                                                                                                 Sequence 40, Application US/10218547 Publication No. US20030100074A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 675; DB 14; Best Local Similarity 100.0%; Pred. No. 3.3e-67; Matches 126; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 39, Application US/10151882
Publication No. US20030059862A1
GENERAL INFORMATION:
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                              FILE REFERENCE: PF561
CURRENT APPLICATION NUMBER: US/10/218,547
CURRENT FILING DATE: 2002-08-15
PRIOR APPLICATION NUMBER: 60/312,542
PRIOR FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 60/330,761
PRIOR APPLICATION NUMBER: 60/330,761
PRIOR FILING DATE: 2001-10-30
                                                                                                                                                          APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Methods And Compositions
TITLE OF INVENTION: Human Endokine Alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: PF554
CURRENT APPLICATION NUMBER: US/10/151,882
CURRENT FILING DATE: 2002-05-22
PRIOR APPLICATION NUMBER: 60/293,100
PRIOR FILING DATE: 2001-05-24
                    NUMBER OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Antibodies Against Tumor Necrosis Factor Delta (APRIL)
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 SEQ ID NOS:
PatentIn ve
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version
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Pred. No. 3.3e-67;
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; TYPE: PRT
; ORGANISM: human
US-10-310-793-44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: 60/336,695
PRIOR TILING DATE: 2001-12-07
PRIOR FILING DATE: 2001-12-07
PRIOR PELICATION NUMBER: 10/226,294
PRIOR FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: 60/314,381
PRIOR FILING DATE: 2001-08-24
PRIOR PELICATION NUMBER: 09/899,059
PRIOR PILING DATE: 2001-07-06
PRIOR PILING DATE: 2001-07-06
PRIOR APPLICATION NUMBER: 60/278,449
PRIOR APPLICATION NUMBER: 60/216,879
PRIOR APPLICATION NUMBER: 60/216,879
PRIOR APPLICATION NUMBER: 60/216,879
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                                                                       Matches
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Publication No. US20030198640A1
                                                                                                               Query Match
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Best Local Similarity
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                                                                                                                                                                                                                                                                       Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 71
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1999-05-13
PRIOR APPLICATION NUMBER: 60/132,227
PRIOR FILING DATE: 1999-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-07-07
PRIOR APPLICATION NUMBER: 09/559,290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methods and Compositions For Treating Inflammatory Bowel Diseases TITLE OF INVENTION: Relating To Human Tumor Necrosis Factor-Gamma Beta FILE REFERENCE: PF573
CURRENT APPLICATION NUMBER: US/10/310,793
CURRENT FILING DATE: 2002-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Yu, Guo-Liang
APPLICANT: Ni, Jian
APPLICANT: Rosen, Craig
                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/134,067
                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/180,908
PRIOR FILING DATE: 2000-02-08
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1 ETAKEPCMAKEGPLPSKMQMASSEPPCVNKVSDWXLEILQNGLYLIYGQVAPNANYNDVA
                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhang, Jun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rosen, Craig A
                                                                100.0%; Score 675; DB 14; ilarity 100.0%; Pred. No. 3.3e-67; Conservative 0; Mismatches 0;
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                                                                    Indels
                                                                                                          Length
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                                          US-10-338-083-17
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APPLICANT: Human Genome Sciences, Inc.,
TITLE OF INVENTION: Heteromultimeric TNF Ligand Family members
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 40, Application US/10202062 Publication No. US20040038349A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.0 SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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Query Match
                                                                                                                                    SOFTWARE: PatentIn version 3.2 SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/307,838
PRIOR FILING DATE: 2001-07-27
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                 APPLICANT: Desjarlais, John R.
APPLICANT: Tansey, Malu Lourdes G.
APPLICANT: Tansey, Malu Lourdes G.
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: Dominant Negative Proteins and Methods Thereof
FILE REFERENCE: A-71273-2
CURRENT APPLICATION NUMBER: US/10/338,083
CURRENT FILLING DATE: 2003-01-16
PRIOR APPLICATION NUMBER: US 60/345,805
PRIOR FILING DATE: 2002-01-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: PF559
CURRENT APPLICATION NUMBER: US/10/202,062
CURRENT FILING DATE: 2002-07-25
                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR FILING DATE: 2002-04-17
                                                                  LENGTH: 113
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: human
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nes 126; Conservative
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    90.1%;
    Score 608;
       DB 14;
       Length 113;
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APPLICANT: Yamamoto, Robert T.

APPLICANT: X1, H. Howard

TITLE OF INVENTION: Identification of Essential Genes in TITLE OF INVENTION: Prokaryotes

FILE REFERENCE: ELITRA.011A

CURRENT APPLICATION NUMBER: US/09/815,242

CURRENT FILING DATE: 2001-03-21

PRIOR APPLICATION NUMBER: 60/191,078

PRIOR APPLICATION NUMBER: 60/206,848

PRIOR FILING DATE: 2000-03-23

PRIOR FILING DATE: 2000-05-23
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US-10-611-399-17
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-611-399-17
                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-815-242-12610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.2 SEQ ID NO 17
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                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12610, Application US/09815242 Patent No. US20020061569A1
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PRIOR PILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: US 60/373,453
PRIOR FILING DATE: 2002-04-17
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2003-07-01
PRIOR APPLICATION NUMBER: US 10/338,083
PRIOR FILING DATE: 2003-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Desjariais, John R.
APPLICANT: Tansey, Malu Lourdes G.
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: DOMINANT NEGATIVE PROTEINS
                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: A-71273-3
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Haselbeck, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                113;
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                                                                                                                                                                                                                                         Carr, Grant J.
                                                                                                                                                                                                                                                                Wall, Daniel
Trawick, John D.
                                                                                                                                                                                                                                                                                    Ohlsen, Kari L.
Zyskind, Judith W.
Wall, Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90.1%; Score 608; DB 16; 100.0%; Pred. No. 5.7e-60; tive 0; Mismatches 0;
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RESULT 12
US-09-757-041-2
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Patent No. US20020009726A1
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PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/269,308 PRIOR FILING DATE: 2001-02-16
                                                                                                       TELEFAX: (619) 535-89. INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                      CLASSIFICATION: PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Reed, John C.
APPLICANT: Sato, Takaaki
TITLE OF INVENTION: CD40 Associated Proteins
NUMBER OF SEQUENCES: 17
MOLECULE TYPE:
                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Campbell, Cathryn
                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                         TELEPHONE: (619) 535-9001
                                                                                                                                                                               NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Diego
STATE: California
                                    TOPOLOGY:
                                                   TYPE: amino acid
                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                    LENGTH:
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FILING DATE: 2000-05-26
APPLICATION NUMBER: 60/242,578
FILING DATE: 2000-10-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                74 QTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLANPQFIS 126
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Similarity 27.4%;
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SEQ ID NOS: 14110
                                                                      543 amino acids
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ID NO: 2:
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Pred. No. 3.5e+02;
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RESULT 13
US-10-004-378A-35
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Best Local
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                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/004,378A CURRENT FILING DATE: 2001-10-24 PRIOR APPLICATION NUMBER: 60/242,882 PRIOR FILING DATE: 2000-10-24 PRIOR APPLICATION NUMBER: 60/242,765 PRIOR FILING DATE: 2000-10-24 PRIOR FILING DATE: 2000-10-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                             PRIOR APPLICATION NUMBER: 60/243,591
PRIOR FILING DATE: 2000-10-26
PRIOR APPLICATION NUMBER: 60/243,950
PRIOR FILING DATE: 2000-10-27
                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/300,206
PRIOR FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/242,789
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                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: No. US20030228301A1el Human TITLE OF INVENTION: Methods of Using the Same
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                                                                                                                                                              OR APPLICATION NUMBER: 60/242,789
OR FILING DATE: 2000-10-24
OR APPLICATION NUMBER: 60/242,768
OR FILING DATE: 2000-10-24
OR APPLICATION NUMBER: 60/242,767
OR FILING DATE: 2000-10-24
OR APPLICATION NUMBER: 60/243,622
                                                                                                            FILING DATE: 2000-10-26
APPLICATION NUMBER: 60/273,047
FILING DATE: 2001-03-02
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               Prior Application SEQ ID NOS: 191
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PatentIn Ver.
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Gangolli, I
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Gunther, Erik
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Grosse, William M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gerlach, Valerie
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Tchernev, Velizar 1
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Spytek, Kimberly A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alsobrook II, John
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28.3%;
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                               See
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                             File Wrapper or PALM
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US-08-813-323A-1
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LENGTH: 543
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08813323A Publication No. US20020031522A1 GENERAL INFORMATION:
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                                                                                  Matches
                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 278-0400
TELEFAX: (212) 391-0525
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION UNMBER: 28,678
REFERENCE/DOCKET NUMBER: 5065
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/813,323A
                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 566 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ye, Zheng-sheng
TITLE OF INVENTION: TRUNCATED CRAF1 INHIBITS CD40 SIGNALING
                                                                                                                                                                                                          NAME/KEY:
                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                               TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Cooper & Dunham, LLP STREET: 1185 Avenue of the Americas
                                                                                                                                                                                       LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                282 EMLR--NNESKI------LHLQRVID----SQAEKLK 306
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252 ASSAVQHVNLLKEWSNSLEKKVSLLQN-----ESVEKNKSIQSLHNQICSFEIEIERQK 305
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                                     ASSEPPCVNKVSDW-----KLEILQNGLYLIYGQVAPNAN----YNDVAPFEVRLYKNK 70
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Cleary, Aileen
                                                                            11.0%; Score 74; DB 8; Length 566; ilarity 28.3%; Pred. No. 25; Conservative 17; Mismatches 24; Indels
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Search completed: November Job time: 127 secs
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US-10-242-212-7
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                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.0 SEQ ID NO 7
                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: AHUJA, SERMA
APPLICANT: BONEWALD, YUNDA
TITLE OF INVENTION: CD40 LIGAND AND CD40 AGONIST COMPOSITIONS AND
TITLE OF INVENTION: METHODS OF USE
FILE REPERENCE: 4003.001000
CURRENT APPLICATION NUMBER: US/10/242,212
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US/09/645,926
PRIOR PELICATION NUMBER: US/09/645,926
PRIOR PELICATION NUMBER: US/09/645,926
PRIOR FILING DATE: 2000-08-24
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 567
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                                                                                    306 EMLR--NNESKI----
                                                                                                                                                                      252 ASSAVQHVNLLKEWSNSLEKKVSLLQN-----ESVEKNKSIQSLHNQICSFEIEIERQK 305
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                                                                                                                            71 DMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLK 109
                                                                                                                                                                                                                21 ASSEPPCVNKVSDW-----KLEILQNGLYLIYGQVAPNAN----YNDVAPFEVRLYKNK 70
                                                                                                                                                                                                                                                          28;
                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                            11.0%; Score 74; DB 28.3%; Pred. No. 25;
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                   2004, 13:10:30
                                                                                                                                                                                                                                                        17; Mismatches
                                                                                    --LHLQRVID----SQAEKLK 330
                                                                                                                                                                                                                                                                                                DB 14;
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Result No.

Query Match

Length

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Description

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Minimum
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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DB
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seq length: 2000000000
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     6730630 seqs, 1107998698 residues
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1 ETAKEPCMAKFGPLPSKWQM.....
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                                                                                                                                                                                                                                                                                                                                                                            November 2,
                                                                                                                                                                                                                                   Pending_Patents_AA Main: *
ETAKEPCMAKFGPLPSKWQM.....VLKNNTYWGIILLANPQFIS
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 1 PCT-US00-04722-2 Sequence 2, Application PC/TU; GENERAL INFORMATION: APPLICANT: Human Genome S: APPLICANT: Rosen, Craig A TITLE OF INVENTION: Human CORRESPONDENCE ADDRESS: DODRESSEE: STERNE, KESS: STREET: 1100 NEW YORK A CITY: WASHINGTON STATE: DC COMDUTER READABLE FORM: MEDIUM TYPE: Floppy disi COMPUTER: IBM PC compat:	1 675 100.0 169 2 675 100.0 169 3 675 100.0 169 4 675 100.0 177 6 675 100.0 177 7 675 100.0 177 7 675 100.0 177 10 675 100.0 177 11 675 100.0 177 12 675 100.0 177 13 675 100.0 177 14 675 100.0 177 15 675 100.0 177 16 675 100.0 177 17 675 100.0 177 18 675 100.0 177 19 675 100.0 177 19 675 100.0 177 20 675 100.0 177 21 675 100.0 177 22 675 100.0 177 23 608 90.1 13 24 608 90.1 17 25 676 83.9 90.1 13 26 444 65.8 90.1 17 27 678 100.0 177 28 648 59.1 103 29 84.5 12.5 1221 30 84.5 12.5 1221 31 77 11.4 217 35 77 11.4 217 36 77 11.4 217 36 77 11.4 217 37 75.5 11.2 3829 44 75.5 11.2 3829 44 75.5 11.2 3829 45 75.5 11.2 5755	
S0004 cienc Endo LER, VE., VE.,	ALIGNMENTS	1 PCT-USO0-04722-2 27 US-10-136-511-2 1 PCT-US96-13282-2 27 US-10-136-551-2 1 PCT-US91-46625-3 1 PCT-US01-46625-3 1 PCT-US02-23782-40 1 PCT-US02-23782-40 1 PCT-US02-23782-16 20 US-09-613-972-14 20 US-09-613-972-14 20 US-09-613-972-14 20 US-09-613-972-14 20 US-10-116-378-16 27 US-10-116-378-16 28 US-10-202-662-40 28 US-10-218-547-40 29 US-10-310-793-44 30 US-10-405-027-3007 30 US-10-435-032-9 30 US-10-495-027-3007 31 US-10-831-4778 32 US-10-831-4778 33 US-10-831-4778 34 US-10-831-4778 35 US-09-307-140-897 36 US-10-485-032-2 37 US-09-813-996A-54421 38 US-09-513-996A-54421 39 US-09-513-996A-54421 39 US-09-513-996A-54421 39 US-09-913-996A-54421 30 US-09-913-913-913-913-913-913-913-913-913-91
s of Use		Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3, Appli Sequence 3, Appli Sequence 40, Appl Sequence 40, Appl Sequence 11, Appli Sequence 14, Appli Sequence 14, Appli Sequence 14, Appli Sequence 14, Appli Sequence 16, Appli Sequence 17, Appli Sequence 40, Appli Sequence 40, Appli Sequence 40, Appli Sequence 40, Appli Sequence 5, Appli Sequence 7, Appli Sequence 17, Appli Sequence 17, Appli Sequence 17, Appli Sequence 17, Appli Sequence 47785, A Sequence 17, Appli Sequence 20671, Appli Sequence 353, Appli Sequence 1042, Appli Sequence 20671, Appli Sequence 20671, Appli Sequence 20671, Appli Sequence 36422, Appli Sequence 36421, Appli Sequence 36421, Appli Sequence 3667, Appli Sequence 3671, A

SOFTWARE:

PatentIn Release #1.0, Version #1.30

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCT-US96-13282-2
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Best Local Similarity
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APPLICATION NUMBER: US 60
FILING DATE: 28-WAY-1999
ATTORNEY/AGENT INFORMATION:
                                                     COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OFFWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/13282
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LENGTH: 169 amino acids
                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C
STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Human Endokine Alpha
NUMBER OF SEQUENCES: 10
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                   STREET: 1100 NEW CITY: WASHINGTON
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                   CLASSIFICATION:
                                        FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Rockville, MD 20850
United States of America
APPLICANTS/INVENTORS: Yu, Guo-Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Genome Sciences, Inc.
9410 Key West Avenue
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/513,584
PRIOR FILING DATE: 2000-02-25
PRIOR PELICATION NUMBER: 09/345,790
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 60/136,788
PRIOR FILING DATE: 1999-05-28
PRIOR APPLICATION NUMBER: 60/122,099
PRIOR APPLICATION NUMBER: 60/122,099
PRIOR PILING DATE: 1999-05-26
PRIOR APPLICATION NUMBER: 60/912,227
PRIOR APPLICATION NUMBER: 60/024,058
PRIOR APPLICATION NUMBER: 60/024,058
                                                                                                                                                                                                                                                                                                                SEQ ID NO 2
LENGTH: 169
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 169 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rosen, Craig A.
TITLE OF INVENTION: Human Endokine Alpha and Methods of Use
FILE REFERENCE: 1488.0470007/EKS/PSC
CURRENT APPLICATION NUMBER: US/10/136,511
CURRENT FILING DATE: 2002-05-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Yu, Guo-Liang APPLICANT: Ni, Jian
                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1996-08-16
NUMBER OF SEQ ID NOS: 10
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REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
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104 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
                                                                                                                                                                     126; Conservative
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                        61 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
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                                                                               44 ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
                                                                                                            1 ETAKEPCMAKFGPLPSKWQMASSSPPCVNKVSDWKLEILQNGLYLIYGQVAFNANYNDVA 60
                                                                                                                                                                 h 100.0%; Score 675; DB 27; Similarity 100.0%; Pred. No. 8.3e-71; O; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : 169 amino acids
amino acid
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GENERAL INFORMATION:
APPLICANT: Genaissance Pharmaceuticals, Inc.
APPLICANT: Bentivegna, Steven C
APPLICANT: Bieglecki, Karyn M
APPLICANT: Sun, Xiao
TITLE OF INVENTION: HAPLOTYPES OF THE TNESF18 GENE
                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 39, Application PC/TUS0216106
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Antibodies Against Tumor Necrosis Factor Delta (APRIL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 126;
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
                                                                                                                                                                                                                                                                   SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                     FILE REFERENCE: PF554PCT
CURRENT APPLICATION NUMBER: PCT/USO2/16106
CURRENT FILING DATE: 2002-05-22
PRIOR APPLICATION NUMBER: 60/293,100
PRIOR FILING DATE: 2001-05-24
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin version 3.0
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CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 60/247,501
PRIOR FILING DATE: 2000-11-10
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                             ORGANISM: Homo sapiens
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                                                                                                                Conservative
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                                                                                                                                100.0%; Score 675; DB 1; 100.0%; Pred. No. 8.8e-71;
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SEQ ID NO 40
SEQ IT NO 40
TYPE: PRT
TYPE: PRT
ORGANISM: human
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 40
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                                                                         Matches 126;
                                                                                          Query Match
Best Local (
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CURRENT FILING DATE: 2002-07-25
PRIOR APPLICATION NUMBER: 60/307,838
PRIOR FILING DATE: 2001-07-27
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Methods And Compositions For Treating Metabolic Bone Diseases Rel
TITLE OF INVENTION: Human Endokine Alpha
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                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PF561PCT
                                                                                                                                                                TYPE: PRT
ORGANISM: human
                                                                                                                                                                                                  LENGTH: 177
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                 1 ETAKEPCMAKPGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA 60
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 ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
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                                                                       100.0%; Score 675; DB 1; 100.0%; Pred. No. 8.8e-71; rative 0; Mismatches 0;
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; Sequence 16, Application US/09247225A
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Goddard, Audrey
; APPLICANT: Gurney, Austin
; APPLICANT: Marsters, Scot A.
; APPLICANT: Pitti, Robert M.
; APPLICANT: Wood, William
; APPLICANT: Wood, William
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GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Mark, Melanie R.
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NUMBER OF SEQ ID NOS: 31
SEQ ID NO 16
LENGTH: 177
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SEQ ID NO 1
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CURRENT APPLICATION NUMBER: US/09/195,368A
CURRENT FILING DATE: 1998-11-18
EARLIER APPLICATION NUMBER: US 60/069,661
EARLIER FILING DATE: 1997-12-12
EARLIER FILING DATE: 1997-11-18
EARLIER FILING DATE: 1997-11-18
                                                                            CURRENT APPLICATION NUMBER: US/09/247,225A CURRENT FILING DATE: 1999-02-09 EARLIER APPLICATION NUMBER: US 60/074,087 EARLIER FILING DATE: 1998-02-09
                                                                                                                                                                                          TITLE OF INVENTION: NOVEL TUMOR NECROSIS FACTOR RECEPTOR HOMOLOG AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P1206R1
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US-09-613-972-14
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US-09-247-225-16
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SEQ ID NO 14
LENGTH: 177
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APPLICANT: P. Mickey Williams
APPLICANT: Mary E. Gerritsen
TITLE OF INVENTION: PROMOTION OR INHIBITION OF ANGIOGENESIS AND
TITLE OF INVENTION: CARDIOVASCULARIZATION BY TUMOR NECROSIS FACTOR
TITLE OF INVENTION: LIGAND/RECEPTOR HOMOLOGS
FILE REFERENCE: P1765R1
CURRENT APPLICATION NUMBER: US/09/613,972A
CURRENT FILING DATE: 2000-07-11
PRIOR APPLICATION NUMBER: US 60/143,304
PRIOR FILING DATE: 1999-07-12
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APPLICANT: Mary E. Gerritsen
TITLE OF INVENTION: PROMOTION OR INHIBITION OF ANGIOGENESIS AND
TITLE OF INVENTION: CARDIOVASCULARIZATION BY TUMOR NECROSIS FACTOR
TITLE OF INVENTION: LIGAND/RECEPTOR HOMOLOGS
FILE REFERENCE: P176SR1
CURRENT APPLICATION NUMBER: US/09/613,972
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PRIOR APPLICATION NUMBER: US 60/143,304
PRIOR FILING DATE: 1999-07-12
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US-10-080-455-1
; Sequence 1, Application US/10080455; GENERAL INFORMATION:
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Gurney, Austin L.
                           RESULT 13
US-10-116-378-16
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                                                                                                                                                                                                                                                                                                                           US-10-080-455-1
                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/080,455;
CURRENT FILING DATE: 2002-02-22
PRIOR APPLICATION NUMBER: 09/195,368
PRIOR FILING DATE: 1998-11-18
PRIOR APPLICATION NUMBER: US 60/069,661
PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: US 60/065,635
PRIOR FILING DATE: 1997-11-18
NUMBER OF SEQ ID NOS: 8
Sequence 16, Application US/10116378 GENERAL INFORMATION:
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SEQ ID NO 14
LENGTH: 177
                                                                                                                                                                                                                                                                 Matches 126;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Marsters, Scot A.
APPLICANT: Pitti, Robert M.
TITLE OF INVENTION: DNA19355 Polypeptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: P1150R2
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 177
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                                                                                                                                                                                                       ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVA
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Conservative 0;
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RESULT 14
US-10-151-882-39
US-10-151-882-39, Application US/10151882
; GENERAL INFORMATION:
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SEQ ID NO 16
LENGTH: 177
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SEQ ID NO 39
LENGTH: 177
                                                                                                                                                                                                                            Query Match 100.0%; Score 675; DB 27; Best Local Similarity 100.0%; Pred. No. 8.8e-71; Matches 126; Conservative 0; Mismatches 0;
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TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P1206R1
CURRENT APPLICATION NUMBER: US/10/116,378
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION TO THE PRIOR APPLICATION NUMBER: US/10/116,378
                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/151,882
CURRENT FILING DATE: 2002-05-22
PRIOR APPLICATION NUMBER: 60/293,100
PRIOR FILING DATE: 2001-05-24
NUMBER OF SEQ ID NOS: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Antibodies Against Tumor Necrosis Factor Delta (APRIL)
FILE REFERENCE: PF554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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PRIOR FILING DATE: EARLIER FILING DATE: 199-02-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/074,087
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: NOVEL TUMOR NECROSIS FACTOR RECEPTOR HOMOLOG
TITLE OF INVENTION: NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Marsters, Scot A
APPLICANT: Pitti, Robert M.
APPLICANT: Wood, William
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                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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121 NPQFIS
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                                                                                      61 PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
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                                                            PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
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Gurney, Austin
Marsters, Scot A.
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172 NPQFIS 177

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RESULT 15

US-10-170-205E-28226

JS-10-170-205E-28226

Application US/10170205E

GENERAL INFORMATION:

APPLICANT: ADAMS, Mark

TITLE OF INVENTION: DEVICES, SUCH AS ARRAYS, COMPRISED OF HUMAN PROTEINS OR PROTEIN

TITLE OF INVENTION: CAPTURE AGENTS, AND USES THEREOF

FILE REFERENCE: CL001381

CURRENT APPLICATION NUMBER: US/10/170,205E

CURRENT APPLICATION NUMBER: 2002-06-13

NUMBER OF SEQ ID NOS: 40312

SOFTWARE: Patentin version 3.2

SEG ID NO 28226
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Search completed: November 2, Job time: 462 secs
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-170-205E-28226
                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 675; DB 27; Length 177; Best Local Similarity 100.0%; Pred. No. 8.8e-71; Matches 126; Conservative 0; Mismatches 0; Indels 0;
                                                                       172 NPOFIS 177
                                                                                                                                                 61 PFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA 120
                                                                                                             121 NPQFIS 126
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OM protein - protein search, using sw model

Run on:

November 2, 2004, 12:55:37; Search time 12 Seconds
(without alignments)
57.744 Million cell updates/sec
Title:
Perfect score: 675
Sequence:
1 ETAKEPCMAKFGPLPSKWQM......VLKNNTYWGIILLANPQFIS 126
Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
Searched:
40726 seqs, 5499444 residues
Total number of hits satisfying chosen parameters:
40726 maximum DB seq length:
Maximum Match 100*
Maximum DB seq length: 200000000

Post-processing: Minimum Match 100*
Listing first 45 summaries

Pending Patents AA New:*
1: /cgn2_6/ptodata/1/paa/USO7_NEW_COMB.pep:*
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5: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*
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8: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*
8: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*
8: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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62.5	63	63	63	63	63	63	63	63.5	63.5	64	64	64	64	64	66.5	67	67	67	67	67.5	68	70.5	608	675	Score
9.3	9.3		9.3				9.3		9.4			9.5	9.5							10.0	10.1	10.4	90.1	100.0	Query
322	233	157	157	157	157	150	147	259	141	307	307	307	157	157	525	285	285	285	285	317	840	137	113	177	Length
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US-10-948-576-138	PCT-US04-31524-147	7-	US-10-957-543-1	US-10-957-134-1	US-10-668-178-1	US-10-959-537-28	US-10-794-751-1	US-60-613-702-11	US-10-794-751-8	US-10-948-576-135	US-10-948-576-123	US-10-948-576-122	US-10-668-178-16	US-10-668-178-15	US-10-076-157A-11	US-10-626-914-9	US-10-964-241-24	US-10-157-779-24	US-10-955-952-24	US-10-948-576-145	US-10-952-698-181	US-10-220-366A-15820	US-10-794-751-17	US-10-959-537-16	ID
138,	147	e 1,	e 1,	e 1,	1, 2	e 28,	Sequence 1, Appli	e 11	е 8,	13	e 12	e 122,	`	,_	e 11,	Sequence 9, Appli	24	24,	24, 7	Sequence 145, App	e 181,	Sequence 15820, A	Sequence 17, Appl	Sequence 16, Appl	Description

RESULT US-10-7 Seque ; Seque ; GENER	QY Db	¥ a b o	RESULT 1 US-10-959-1 Sequence GENERAL 1 APPLICAL		00000000000000000000000000000000000000
ESULT 2 S-10-794-751-17 Sequence 17, Applicat GENERAL INFORMATION: APPLICANT: Desjarlai	1 ETAK 52 ETAK 61 PFEV 112 PFEV 111 NPQF 172 NPQF	H: 177 PRT ISM: Hom -537-16 atch cal Sim: 126;	116, 1 116, 1 116, 1 117, 1 117, 1 117, 1 117, 1 1 117, 1 1 117, 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		и
plicati TION: jarlais	TAKEPCMAKFG TAKEPCMAKFG TAKEPCMAKFG FEVRLYKNKDM FEVRLYKNKDM FEVRLYKNKDM FEVRLYKNKDM FOFIS 126	no sapiens ilarity Conservat	7-16 6, Application US FORMATION: 1. Assikenazi, Avi 2. Goddard, Audre 2. Gurney, Austin 2. Goddard, Scot 2. Pitti, Robert 3. Wood, William 1. NOVEL 4. NOVEL 5. WOOD, WILLIAM 1. NOVEL 5. WOOD, WILLIAM 1. NOVEL 5. WOOD 6. PITTION NOVE 6. PITTION NUMBER 7. LING DATE: 1999-02 7. SEQ ID NOS: 31 7. SEQ ID NOS: 31		
on US/1 , John	MIQO HIQO	100.0 100.0	ion US/10959537 Audrey Austrin , Scot A. Cobert M. Iliam NOVEL TUMOR NEC ACIDS ENCODING 16R1 2004-10-06 6MBER: US/10/9/247 999-02-09 MBER: US 60/074 998-02-09		1157 6 1122 6 6 2214 6 6 2214 6 6 327 6 6 322 6 6 322 6 6 322 6 6 322 6 7 6 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
.0794751 R.	SEPPCYNKVSDWKLEILQNG 	<pre>\$; Score 675; DB 6; \$; Pred. No. 1.4e-69; 0; Mismatches 0;</pre>	9537 NECROSIS FACTOR DING THE SAME 10/959,537 16 1/247,225	ALIGNMENTS	US-10-668-178-2 US-10-970-713-286 US-10-970-713-306 US-10-948-576-114 US-10-948-576-112 US-10-948-576-129 US-10-948-576-139 US-10-948-576-139 US-10-948-576-139 US-10-948-576-139 US-10-948-576-141 US-10-948-576-141 US-10-948-576-141 US-10-948-576-121 US-10-668-178-3
	LYLIYGQVAENANYNDVA 60 	Length 177; Indels 0; Gaps 0;	RECEPTOR HOMOLOG AND NUCLEIC		Sequence 2, Appli Sequence 286, App Sequence 306, App Sequence 144, App Sequence 142, App Sequence 129, App Sequence 139, App Sequence 139, App Sequence 157, App Sequence 157, App Sequence 2, Appli Sequence 2, Appli Sequence 4, Appli Sequence 5, Appli Sequence 5, Appli Sequence 9, Appli Sequence 9, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 1121, App

Thomason, Adam Read

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Sequence 15820, Application US/10220366A

GENERAL INFORMATION:
APPLICANT: HYSEQ, INC
TITLE OF INVENTION:
CURRENT APPLICATION NOVEL Nucleic Acids and Polypeptides
FILE REFERENCE: 21272-042
CURRENT APPLICATION NUMBER: US/10/220,366A
CURRENT FILING DATE: 2002-08-28
PRIOR APPLICATION NUMBER: 09/577,409
PRIOR APPLICATION NUMBER: 09/577,409
PRIOR FILING DATE: 2000-05-18
PRIOR APPLICATION NUMBER: 09/515,126
PRIOR FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 27802
SOFTMARE: CITATUM
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                                                                                                                                                                                                                                                       ; NAME/KEY: misc feature
; LOCATION: (1)...(137)
; OTHER INFORMATION: Xaa = any amino acid or nothing
US-10-220-366A-15820
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                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 15820
LENGTH: 137
                                                                                                                                                                            Matches
                                                                                                                                                                                                                Query Match
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Best Local Similarity
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CURRENT FILING DATE: 2004-03-05
PRIOR APPLICATION NUMBER: US 10/338,083
PRIOR FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/452,707
PRIOR FILING DATE: 2003-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Custom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/482,081
PRIOR FILING DATE: 2003-06-23
PRIOR APPLICATION NUMBER: US 60/523,880
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: A-72175-1
                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                                                                                                                                                                       / Match 10.4%;
Local Similarity 30.0%;
nes 30; Conservative 1
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                                      GDTIDLIENSEHQVLKNNTYW------
                                                                                 LSIDQVG---VHRSIAANMTFAEIVTPF-----NIDRLQELVRRGNSQYPGAKYIIRDN 97
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GDRIDLRFHPKHSDLHLQTGYKVKRHMCDGDIVIFNRQCI 137
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                                                                                                                                                                          13;
                                                                                                                                                                       Score 70.5; DB Pred. No. 0.25; 3; Mismatches
                                      GIILLANPQFI 125
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US-10-948-576-145
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US-10-952-698-181
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CURRENT FILING DATE: 2004-09-28
NUMBER OF SEQ ID NOS: 207
SOFTWARE: Patentin version 3.2
SEQ ID NO 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 145,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 181, Application: GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mack, David APPLICANT: Aziz, Nata
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quence 145, Application US/10948576 GENERAL INFORMATION:
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              ZIP: 75201

COMPUTER READABLE FORM:

MEDIUM TYPE: CD

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/948,576
FILING DATE: 22-Sep-2004
CLASSIFICATION: CURKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Carol M
                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  117 -- ILLANPOFIS 126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48 CEAKGIKMVSEISVPPSRP-----FQLSLLNNGLTMLHTNDFSGLTNAISIHLGFNN 99
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                                                                                                                                                                                                                                 STATE: Texas
COUNTRY: USA
                                                                                                                                                                                                                                                                                      ADDRESSEE: Carol M. Nielsen, STREET: P.O. Box 50784
                                                                                                                                                                                                                                                                      CITY: Dallas
APPLICATION NUMBER: 10/083,446
                                                                                                                                                                                                                                                                                                                                                                     OF INVENTION: Methods Of Ex-Vivo Using Multivariant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10952698
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ilarity 18.2%;
Conservative 3
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Paik, Kumnan
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Klein, Barbara K.
McKearn, John P.
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Caparon, Maire H.
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Pred. No. 5.2;
31; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                     Expansion Of Hematopoietic IL-3 Hematopoiesis Chimera
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Best Local (
                         PRIOR FILING DATE: 2002-04-12
PRIOR APPLICATION NUMBER: 60/049911
PRIOR PILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR PILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
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APPLICANT:
APPLICANT:
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                                                                                                                              CURRENT APPLICATION NUMBER: US/10/955,952
CURRENT FILING DATE: 2004-09-29
ERIOR APPLICATION NUMBER: US/10/121,058
                                                                                                                                                                                                  TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                       FILE REFERENCE: P3330R1C12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/446,872
PILING DATE: 06-UUW-1995
ATTORNEY/AGENT INFORMATION:
NAME: Carol M. Nielsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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TELEPHONE: 713-650-2722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        123 GGGSGGSNMAPVPPGE----DSKDVAAPHRQPLTSSERI 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24, Application US/10955952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53 -----QTLTNKSKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 EPCMAKEGPLPSKWQMASSEPPCVNKVSDWK-----LEILQNGLYLIYGQVAP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 37,0 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 22-FEB-2002
APPLICATION NUMBER: 08/7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 317 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                                                                                                                                                      Watanabe, Colin K
Wood, William
                                                                                                                                                                                                                                                                                                                                                        Godowski, Paul J.
Gurney, Austin L.
                                                                                                                                                                                                                                                                                         Tumas, Daniel
                                                                                                                                                                                                                                                                                                                                                                                                            Gerritsen, Mary E.
                                                                                                                                                                                                                                                                                                                                                                                                                              Gao, Wei-Qiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DeForge,Laura
                                                                                                                                                                                                                                                                                                         Stewart, Timothy A
                                                                                                                                                                                                                                                                                                                                           Sherwood, Steven
                                                                                                                                                                                                                                                                                                                                                                                            Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                                                                                                                                Filvaroff, Ellen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Beresini, Maureen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
NUMBER: 60/059115
                1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -LPSA-TAAPSRHPIIIKAGDWQEFREKLTFYLVTLEQAQEQQYVIEGRISP
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14-FEB-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 67.5;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
                                                                                                                                                                  APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRA
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C426
CURRENT APPLICATION NUMBER: US/10/157,779
CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR APPLICATION NUMBER: 60/059352
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR FILING DATE: 1997-09-19
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR TILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059122
PRIOR APPLICATION NUMBER: 60/059122
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NUMBER OF SEQ ID NOS: 550
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ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         247 SAGIAKLEEGDELQLAIPRENAQISLDGDVTFFGALKL 284
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                                                                                                                                                                                                                                                                                        T: Zhang, Zemin
INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       42 GLYLIYGQVAPNANYNDVAPFEVRLYKNK-----DMIQTLTNKSKIQNV------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GYFFIYGQVL----YTDKTYAMGHLIQRKKVHVFGDELSLVTLFRCIQNMPETLPNNSCY 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Baker, Kevin P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10157779
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Godowski, Paul J.
                                                                                                                                                                                                                                                                                                                                                                            Wood, William
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DeForge, Laura
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                                                                                                    PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
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PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 24
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TYPE: PRT
ORGANISM: Homo 9
3-10-157-779-24
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                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C33

CURRENT APPLICATION NUMBER: US/10/964,241

CURRENT FILING DATE: 2004-10-12
                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PRIORT APPLICATION NUMBER: US/10/123,236 PRIOR FILING DATE: 2002-04-15
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                                                                               APPLICATION NUMBER: 60/059113
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059115
FILING DATE: 1997-09-17
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FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/059352
FILING DATE: 1997-09-19
FILING DATE: 1997-09-17
                     APPLICATION NUMBER: 60/059122
                                          FILING DATE:
                                                            APPLICATION NUMBER: 60/059117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                         Wood, William Zhang, Zemin
                                                                                                                                                                                                                                                                                                                                                                                                                                    Watanabe, Colin K
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Godowski, Paul J.
Gurney, Austin L.
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Filvaroff, Ellen
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Pred. No. 1.6;
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US-10-626-914-9
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                                                                                                                                                                                                                                                                                               US-10-626-914-9
                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9, Application US/10626914 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 24
                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/626,914
CURRENT FILING DATE: 2003-07-25
PRIOR APPLICATION NUMBER: US 60/398,530
PRIOR FILING DATE: 2002-07-25
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: CHUNTHARAPAI, ANAN
APPLICANT: GREWAL, IQBAL
APPLICANT: KIM, KYUNG JIN
APPLICANT: YAN, MINHONG
TITLE OF INVENTION: TACI Antibodies and Uses Thereof
FILE REFERENCE: P1942R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                LENGTH: 285
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    140 ETVTQDCLQ-----LIADSETPTIQKGSYTFVPWLLSFKRGSALEEKENKILVKET 190
                                                                                                                                             140 ETVTQDCLQ-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    247 SAGIAKLEEGDELQLAIPRENAQISLDGDVTFFGALKL 284
                                                                                                           42 GLYLIYGQVAPNANYNDVAPFEVRLYKNK-----DMIQTLTNKSKIQNV--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42 GLYLIYGQVAPNANYNDVAPFEVRLYKNK-----DMIQTLTNKSKIQNV------
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                                                                                                                                                                                                                                     h 9.9%; Score 67; DB 6; Length 285; Similarity 22.2%; Pred. No. 1.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
 SAGIAKLEEGDELQLAIPRENAQISLDGDVTFFGALKL
                                  -GGTYELHVGDTIDLIFNSEHQVLK---NNTYWGIILL 119
                                                                      GYFFIYGQVL----YTDKTYAMGHLIQRKKVHVFGDELSLVTLFRCIQNMPETLPNNSCY 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -GGTYELHVGDTIDLIFNSEHQVLK---NNTYWGIILL 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ETAKEPCMAKFGPLPSKWQMASSEPPCVNKVS----DW------KLEILQN 41
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                                                                                                                                                                                                                    Conservative
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                                                                                                                                                 --LIADSETPTIQKGSYTFVPWLLSFKRGSALEEKENKILVKET 190
                                                                                                                                                                                                                      20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 67;
Pred. No. 1
                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6;
                                                                                                                                                                                                                      51; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 285
 284
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                                                                                                                                                                                                                      52;
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RESULT 10 US-10-076-157A-11

; Sequence 11, Application US/10076157A

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US-10-668-178-15
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; ORGANISM: Ashbya gossypii
US-10-076-157A-11
                                                                                                                                                                        Query Match
Best Local S
Matches 26
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Matches
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15, Application US/10668178
GENERAL INFORMATION:
APPLICANT: KABUSHIKI KAISHA HAYASHIBARA SEIBUTSU KAGAKU KENKYUJO
APPLICANT: MAYUMI, Tadanori
APPLICANT: TSUTSUMI, Yasuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: JP 83509/2002
PRIOR FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: JP 185387/2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: MAYUMI2A CURRENT APPLICATION NUMBER: US/10/668,178 CURRENT FILING DATE: 2003-09-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: TSUTSUMI, Yasuo
APPLICANT: NAKAGAWA, Shinsaku
APPLICANT: IKEGAMI, Hakuo
TITLE OF INVENTION: Biologically-active conjugate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/076,157A
CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: Germany, 19757755.5
PRIOR FILING DATE: 1997-12-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Garcia, Maria Angeles Santos
TITLE OF INVENTION: Phosphoriboxyl-Pyrophosphate Synthetase Polypeptide
FILE REFERENCE: PF48687-2/DP
                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 21
SOFTWARE: WordPerfect 8
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                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                    LENGTH: 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 525
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                    102 QRETPEGAEALPWYEPIYLGGVFQLETGDRLSAEINRPDYLDFAESGQV-----YFGIIA 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99
                                                       81 KIQN------VGGTYELHVGDTID-----LIFNSEHQVLKNNTYWGIIL
                                                                                                                                 30 KVSDWKLEILQNGLYLIYGQV-----APNANY---NDVAPFEVRLYKNKDMIQTLTNKS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26 PCVNKVSD--WKLE--ILQNGLYLIYGQVAPNANYNDVAPFEVRL-----YKNKDMIQT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76 LTNKSKIQNVG-----GTYELHVGDTIDLIFNSEHQVLKNNTYW 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29;
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                                                                                            BLRDNQLVVPSEGLYLIYSQVLFSGQGCPSTHVLLTHTISRIAVSYQTRVNLLSAIASPC
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Hoeffken, Hans Wolfgang
Doval, Jose Luis Revuelta
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                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                2002-06-26
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                                                                                                                                                                                            9.5%;
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16; Mismatches
                                                                                                                                                                        Score 64; DB 6; Length 157; Pred. No. 1.6; Indels 1; Mismatches 38; Indels
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Pred. No. 4.1;
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LENGTH: 157
TYPE: PRT
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APPLICANT: MAYUMI, Tadanori
APPLICANT: TSUTSUMI, Yasuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/668,178
CURRENT FILING DATE: 2003-09-24
PRIOR APPLICATION NUMBER: JP 83509/2002
PRIOR FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: JP 185387/2002
PRIOR APPLICATION NUMBER: JP 185387/2002
PRIOR FILING DATE: 2002-06-26
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TITLE OF INVENTION: Biologically-active conjugate
FILE REFERENCE: MAYUMIZA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                          TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              157 L 157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   81 KIQN-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42 ELRDNQLVVPSEGLYLIYSQVLFSGQGCPSTHVLLTHTISRIAVSYQTRVNLLSAIASPC 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30 KVSDWKLEILQNGLYLIYGQV-----APNANY---NDVAPFEVRLYKNKDMIQTLTNKS 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 9.5%;
Similarity 21.5%;
                                      STATE:
                                                                         STREET: P.O. Box 50784
                STATE: Texas
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MAYUMI, Tadanori
TSUTSUMI, Yasuo
NAKAGAWA, Shinsaku
                                                                                              ADDRESSEE: Carol M. Nielsen, Winstead Sechrest & Minick P.C.
ZIP: 75201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          157
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                                                                                                                                     Thomas, John W.

INVENTION: Methods Of Ex-Vivo Expansion Of Hematopoietic Cells
Using Multivariant IL-3 Hematopoiesis Chimera Proteins
F SEQUENCES: 197
                                                                                                                                                                                                         Caparon, Maire H.
Easton, Alan M.
Klein, Barbara K.
McKearn, John P.
Olins, Peter O.
Paik, Kumnan
                                                       Dallas
                                                                                                                                                                                                                                                                                                                                                                     Abrams, Mark A.
                                                                                                                                                                                                                                                                                                         Braford-Goldberg,
Maire H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21;
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                                                                                                                                                                                                                                                                                                                                  Sarah
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36;

Gaps

UI

156

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TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 122:
US-10-948-576-122
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US-10-948-576-123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 123, Application US/10948576
GENERAL INFORMATION:
APPLICANT: Abrams, Mark A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 214-745-5390 INFORMATION FOR SEQ ID NO: 122:
                                                                                                                                                                      Olins, Peter O.
Paik, Kumnan
Thomas, John W.
TITLE OF INVENTION: Methods Of Ex-Vivo
NUMBER OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Carol M. Nielsen
REGISTRATION NUMBER: 37,676
REFERENCE/DOCKET NUMBER: 279
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-650-2722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/446,872 FILING DATE: 06-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/948,576
FILING DATE: 22-Sep-2004
    COMPUTER READABLE FORM:
                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Carol M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: CD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 EPCMAKFGPLPSKWQMASSEPPCVNKVSDWK------LEILQNGLYLIYGQVAP 52
                      CITY: Dallas
STATE: Texas
COUNTRY: USA
ZIP: 75201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGGSGGGSNMATPLGPASSLPQSFLLKSLEQVRKIQGDGAALQEKLCATYKL 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QPC-----LPSA-TAAPSRHPIIIKAGDWQEFREKLTFYLVTLEQAQEQQYVIEGRISP 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 307 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 10/083,446
FILING DATE: 22-FEB-2002
APPLICATION NUMBER: 08/762,227
FILING DATE: 09-DEC-1996
APPLICATION NUMBER: US 08/192,325
FILING DATE: 14-FEB-1994
                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                1: Abrams, Mark A.
Bauer, S. C.
Braford-Goldberg, Sarah R.
Caparon, Maire H.
Easton, Alan M.
Klein, Barbara K.
McKearn, John P.
Olins, Peter O.
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25.9%;
                                                                                                              arol M. Nielsen, Winstead Sechrest & Minick P.C Box 50784
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Pred. No. 3.
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3.9;
                                                                                                                                                                                               Expansion Of Hematopoietic IL-3 Hematopoiesis Chimera
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Proteins
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US-10-948-576-135
; Sequence 135, Applicat;
; GENERAL INFORMATION:
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APPLICATION NUMBER: 10/083,446
ETLING DATE: 22-FEB-2002
APPLICATION NUMBER: 08/762,227
FILING DATE: 09-DEC-1996
APPLICATION NUMBER: US 08/192,325
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: US 08/446,872
FILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Carol M. Nielsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 Match 9.5%; Score 64; DB 6; Length 307; Local Similarity 25.9%; Pred. No. 3.9; les 29; Conservative 11; Mismatches 40; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/948,576
FILING DATE: 22-Sep-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 123:
COUNTRY: USA
ZIP: 75201
COMPUTER READABLE FORM:
MEDIUM TYPE: CD
                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Carol M. Nielsen,
STREET: P.O. Box 50784
                                                                                                                                                                                                 TITLE OF INVENTION: Methods Of Ex-Vivo Expansion Of Hematopoietic Cells
Using Multivariant IL-3 Hematopoiesis Chimera Prote:
NUMBER OF SEQUENCES: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 713-650-2722
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              123 GGGSGGGSNMATPLGPASSLPQSFLLKSLEQVRKIQGDGAALQEKLCATYKL 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 EPCMAKFGPLPSKWQMASSEPPCVNKVSDWK-----LEILQNGLYLIYGQVAP 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 307 amino acids
TYPE: amino acid
                                                                                       STATE: Texas
                                                                                                            CITY: Dallas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 214-745-5390
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REFERENCE/DOCKET NUMBER: 279
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Paik, Kumnan
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                                                                                                                                                                                                                                                                                                                                                  Easton, Alan M.
Klein, Barbara K.
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Braford-Goldberg,
Caparon, Maire H.
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32;

Gaps

5

Proteins

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Db
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Search completed: November Job time: 13 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/083,446
APPLICATION NUMBER: 08/762,227
APPLICATION NUMBER: 08/762,227
FILLING DATE: 09-DEC-1996
APPLICATION NUMBER: US 08/192,325
FILLING DATE: 09-DEC-1994
APPLICATION NUMBER: US 08/446,872
APPLICATION NUMBER: US 08/446,872
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: US 08/446,872
FILING DATE: 10-CUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: CAROL NUMBER: 37,676
REGISTRATION NUMBER: 37,676
REFERENCE/DOCKET NUMBER: 2790/7 US
TELECOMMUNICATION NUMBER: 2790/7 US
TELECOMMUNICATION INFORMATION:
TELECHONE: 713-650-2722
TELEFAX: 214-745-5390
INFORMATION FOR SEQ ID NO: 135:
SEQUENCE CHARACTERISTICS:
LENGTH: 307 amino acids
TYPE: TOROLOGY: 110-67
                                                                                                                                                                                                                                                                                                                                             Query Match 9.5%; Score 64; DB 6; Length 307; Best Local Similarity 25.9%; Pred. No. 3.9; Matches 29; Conservative 11; Mismatches 40; Indels 32; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMMER: US/10/948,576
FILING DATE: 22-Sep-2004
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                   123 GGGSGGGSNMATPLGPASSLPQSFLLKSLEQVRKIQGDGAALQEKLCATYKL 174
                                                                                                                                                                   53 NANYNDVAPFEVRLYKNKDMIQTLTNKS-----KIQNVG------GTYEL 91
                                                                                                                                                                                                                                70 QPC-----LPSA-TAAPSRHPIIIKAGDWQEFREKLTFYLVTLEQAQEQQYVIEGRISP
                                                                                                                                                                                                                                                                                         5 EPCMAKFGPLPSKWQMASSEPPCVNKVSDWK------LEILQNGLYLIYGQVAP 52
                           2, 2004, 13:08:18
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Post-processing: Minimum Match
Maximum Match
Listing first
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Perfect score:
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                                                                                                                             length: 0
length: 2000000000
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PIR_79:*
1: pir1:*
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(c) 1993 - 2004
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	Score	Query Match	Length	DB.	ID	scription
1	77	11.4	851	2	T51545	protein
2	75.5	11.2	314	N	150811	
ω		٠	3890	N	C89921	othetical prot
4	75		575	N	T28230	dor
5	74	11.0	233	N	S11688	tumor necrosis fac
6	74	11.0	543	N	S68467	CD40 receptor-asso
7	74		567	N	149272	CD40 receptor-asso
œ	74		568	N	A55960	-
9	73.5	•	504	N	T47446	-
10	72	10.7	393	N	D75207	hypothetical prote
11	72	10.7	477	N	S32223	' Р
12	72	10.7	486	N	F35965	μ,
13	72	10.7	525	N	E70125	hypothetical prote
14	72	10.7	609	_	A43458	on proce
15	71.5	10.6	1123	N	A72311	hypothe
16	71.5	10.6	3623	N	T08618	
17	71	10.5	232	ب	S12606	٠
18	71	10.5	473	N	S39404	Lin-
19	71	10.5	486		96	, <u> </u>
20	71	10.5	486		96	llin-
21	71		486		B35965	cillin-
22	71		486		C35965	cillin-
23	71	10.5	679	N	S06000	cillin-
24	71		680		G95194	_
25	71	10.5	685		D98061	a
26	70.5	10.4	297		T28188	
27	70.5	10.4	890		A30481	bacteriocin BCN5 -
28	70.5	10.4	1508		T31098	probable dextransu
29	70	10.4	1010		T33372	hypothetical prote

C;Accession: I508i1
R;Betz, U.A.K.; Mayer, W.E.; Klein, J.
Proc. Natl. Acad. Sci. U.S.A. 91, 11065-11069, 1994
A;Title: Major histocompatibility complex class I genes A;Reference number: I50810; MUID:95062206; PMID:7972010
A;Accession: I50811

of the

coelacanth Latimeria cha

ISO811
MHC class I protein, alpha 1, alpha 2, alpha 3 and transmembrane domains - coelacanth C;Species: Latimeria chalumnae (coelacanth)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 09-Jul-2004

A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-314 <BET>

UNIPROT:Q31417; EMBL:U08034; NID:g549857; PIDN:AAA52346.1; PID:g560

A; Cross-references: Genetics:

444	42	40	38	37	36	S	34	ω ω	32	31	30
6,6,0	68. 	58.	68.5	68.5	68.5	68.5	69.5	69.5	69.5	69.5	69.5
10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.3	10.3	10.3	10.3	10.3
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B64565 E71858	PC4035	T20898	VGXR2S	859322	B72666	B35216	B36329	B71617	A54505	S67649	A25451
conserved hypothet hypothetical prote	cell-cycle-depende tumor necrosis fac	hypothetical prote	glycoprotein VP7 p	hypothetical prote		FP14 protein - fow	рr		serine-repear and	- 1	co.

ALIGNMENTS

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T51545	
receptor	receptor protein kinase-like protein - Arabidopsis thaliana
N; Altern	N;Alternate names: protein F2K13_50 C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: C;Access)0 #cext_
R;Sato,	submitted to the Protein Sequence Database, August 2000
A; Refere	A; Reference number: Z25394
A;Access	A;Accession: T51545
A:Molecu	A; Scarus: pretrimerary A:Molecule type: DNA
A;Residu	A; Residues: 1-851 <sat></sat>
A; Cross-	A;Cross-references: UNIPROT:Q9LFL1; EMBL:AL391141
A; Experi	A;Experimental source: cultivar Columbia; BAC clone F2KL3
A:Map posit	A; Map position: 5
A; Intron	A;Introns: 23/1; 202/2; 371/1; 428/2; 452/2; 476/2; 531/1; 602/3; 645/1; 66//1; /30/2 A:Note: F2X13 50
C;Superf	C;Superfamily: protein kinase homology
Query Match Best Local (Matches 2)	Query Match 11.4%; Score 77; DB 2; Length 851; Best Local Similarity 28.0%; Pred. No. 19; Matches 28; Conservative 17; Mismatches 27; Indels 28; Gaps 5;
γ	34 WKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLTNKSKIQNVGGT 88
Db	YMIKAVFVYGNYDGYNDYE
40	89 YELHVGDTIDLIENSEHQVLKNNTY 113
Db	

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A;Note: MSVUby
C;Superfamily: variola virus N3L protein
                                                                                                                                                   A;Status: preliminary; translated A;Molecule type: DNA A;Residues: 1-575 <AFO>
                                                                                                                                                                                                                     C;Date: 21-Jan-2000 #sequence_revision 21-Jan-2000 #text_char (;Accession: T28230 R;Afonso, C.L.; Tulman, E.R.; Lu, Z.; Oma, E.; Kutish, G.F.; J. Virol. 73, 533-552, 1999
A;Title: The genome of Melanoplus sanguinipes entomopoxvirus. A;Reference number: Z20484; MUID:99102612; PMID:9847359
                                                                                                                            A; Cross-references: UNIPROT: Q9YW23; EMBL: AF063866; NID: g4049647;
                                                                                                                                                                                                             A; Accession: T28230
                                                                                                                                                                                                                                                                                                                                                     ORF MSV069 probable rifampicin resistance (RIF) protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecute ',r'.

A;Residues: 1-3890 <KUR>
A;Cross-references: UNIFROT:099U53;
A;Cross-references: strain N315
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A;Title: Whole genome sequencing of meticillin-registant Stapylococcus aureus.

**Peference number: A89758; MUID:21311952; PMID:11418146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, C.; Shiba, T.; Hattori, K.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein ebhB [imported] - Staphylococcus aureus (strain N315) C;Species: Staphylococcus aureus C;Species: Staphylococcus aureus C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004 C;Bccession: C89921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Experimental source: strain C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ф
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Introns: 91/1; 183/1; 276/1
C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
     Query Match
Best Local S
Matches 32
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     32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NNTTNDSQSKHV----SIHVGK----ISEDAHPIVLGNT--EKVVVVNPTAVS 1039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QTLTNKSKIQNVGGTYELHVGDTIDL1FNSEHQVLXNNTYWGIILLANPQFIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LPSKWQMASSEEPCVNKVSDWKLEILQNGLYLIYGQVAPNANYNDVAPFEYRLYKNKDMI 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LPSGWTSNLTK----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEVRLYKNKDMIQTLT 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -EVRVYDRPDLERNLT 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KAQIRSKVCPVDEAWDSYFSE-
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                       23.7%;
                                      11.1%;
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     21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -SDNK---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 75.5; DI
Pred. No. 1.8e-
17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9;
                     Score 75; DB
Pred. No. 19;
                                                                                                                                                                                        from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                        entomopoxvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 75.5;
Pred. No. 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GB:BA000018; PID:g13701233; PIDN:BAB42528.1;
     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                21-Jan-2000 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.5; DB 2;
No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -NGSLAITGRVSMNQAFNSDITFKV---SATDNV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -DWKQECIE-GLKKYLLYGK---ETLERKVAP
                                    2
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   56;
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                                    Length 575;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 3890;
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                                                                                                                                                                                                                                                                                                                                                     Heliothis armigera
 26;
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                                                                                                                                PIDN:AAC97814.1;
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 Gaps
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K.;
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CD40 receptor-associated (man)
C;Species: Homo sapiens (man)
C;Date: 17-Uul-1998 #sequence revision 17-Uul-1998
C;Accession: S68467; I53498
R;Sato, T.; Irie, S.; Reed, J.C.
FEBS Lett. 358, 113-118, 1995
A;Title: A novel member of the TRAF family of putated for the TRAF family of
       A; Gene: CAP-1
C; Superfamily:
C; Keywords: hc
                                                                                                                                                                                                                                                                A;Reference number: I53498; MUID:95:
A;Accession: S68467
A;Status: nucleic acid sequence not
                                                                                                                              A;Cross-references: UNIPROT:Q13114; EMBL:L38509; NID:g695357; PIDN:AAA68195.1; PID:g69535A;Experimental source: tissue-type fetal brain
                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-543 <SAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S68467
CD40 receptor-associated protein CAP-1 -
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C;Superfamily: tumor necrosis factor
C;Superfamily: tumor necrosis factor
C;Keywords: glycoprotein; lipoprotein; myristylation; transmembrane
F;19,20/Binding site: myristate (Iys) (covalent) #status predicted
F;81/Binding site: carbohydrate (Ser) (covalent) #status predicted
F;145-177/Disulfide bonds: #status predicted
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C;Species: Felis silvestris catus (domestic cat)
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT
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C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-233 < MCG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: A; Accession: S11688
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A;Title: Gene sequence of feline tumor necrosis factor alpha.
A;Reference number: S11688; MUID:91016860; PMID:2216740
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                                      Superfamily:
                                                                                                         Genetics:
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   118 ELTDNOLKVESDGLYLIYSOVLFTGQGCESTHVLLTHAISRFAVSYQTKVNLLSAI--KS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81 KIQN----
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       homodimer;
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TNF receptor-associated factor (TRAF); R. modimer; signal transduction; zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WOMASSEPPCVN-----KVSDWKLEILQN-GLYLIYGQV-----APNANYNDVAPFE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YQFTSSSSVYENGYLIYKITDSKLEGCNNFGLIITLPEIKGIGSIRYQSNFVYKLIEEFV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IETIDNDSTVNTIIKKSGLELLFEFNRSGSKYSKIIGNNIDLCSFNTGYTADDIIFTSRE 151
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Pred. No. 7.8;
                                                                                                                                                                                                                                                                          shown
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                                                                                                                                                                                                                                                                                                                                                                     transducing proteins binds
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RESULT 7
149272
CD40 receptor-associated factor 1
                                                      A;Cross-references: UNIPROT:Q13114; EMBL:U21092; NID:97260 R;Mosialos, G.; Birkenbach, M.; Yalamanchili, R.; VanArsda Cell 80, 389-399, 1995 A;Title: The Epstein-Barr virus transforming protein LMP1 A;Reference number: A55649; MUID:95163092; PMID:7859281 A;Accession: A55649
                                                                                                                                                                                                                                                          N;Alternate names: CD40-binding protein
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 15-Mar-1996 #sequence revision 15-Mar-1996 #text_change 09-Jul-2004
C;Accession: A55960; A55649; A55135
R;Cheng, G.; Cleary, A.M.; Ye, Z.S.; Hong, D.I.; Lederman, S.; Baltimore, D. Science 267, 1494-1498, 1995
R;Cheng 267, 1494-1498, 1995
A;Title: Involvement of CRAF1, a relative of TRAF, in CD40 signaling.
A;Reference number: A55960; MUID:95184010; PMID:7533327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Cheng, G.; Cleary, A.M.; Ye, Z.S.; Hong, D.I.; Lederman, S.; Baltiv Science 267, 1494-1498, 1995
A;Title: Involvement of CRAF1, a relative of TRAF, in CD40 signaling A;Reference number: A55960; MUID:95184010; PMID:7533327
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C; Superfamily: TNF re
C; Keywords: zinc fing
F; 48-96/Domain: RING
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A;Cross-references: UNIPROT:Q60803; EMBL:U21050; NID:g719292; PIDN:AAC52175.1;
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A;Status: nucleic acid sequenc
A;Molecule type: mRNA
A;Residues: 1-128,'M',130-568
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                                                                                                                                                                                                    A; Molecule type: mRNA
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                                                                                                                                                                                                                                            A; Accession: A55960
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Best Local S
Matches 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily: TNF receptor-associated factor; Keywords: zinc finger
                                                                                                                                                                                 ;Residues: 1-568 <RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLK 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLK 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ASSAVQHVNLLKEWSNSLEKKVSLLQN-----ESVEKNKSIQSLHNQICSFEIEIERQK
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                                          acid sequence not
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28.3%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 74;
Pred. No.
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                                                                                                                                         NID:g726087; PIDN:AAC50112.1; PID; VanArsdale, T.; Ware, C.; Kieff,
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                                                                                                                                             PID:g7260
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C;Keywords: coiled coil; zinc F;49-97/Domain: RING fine...
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A;Residues: 1-128,'M',130-133,135-404,'G',406-568 <HUA>
A;Cross-references: GB:U15637; NID:g595910; PIDN:AAA56753.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Hu, H.M.; O'Rourke, K.; Boguski, M.S.; Dixit, V.M. J. Biol. Chem. 269, 30069-30072, 1994
A;Title: A novel RING finger protein interacts with the A;Reference number: A55135; MUID:95073988; PMID:7527023
                                                                                                                                                                                                                                                                             A;Cross-references: GB:U19260; NID:g675459; PIDN:AAA65732.1; PID:g675460 R;Hu, H.M.; O'Rourke, K.; Boguski M.S.; Dixit, V.M.
                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: A; Accession: T47446
                                                                                                                                                                                                                                                                                                                                                                                                              R;Jordan, N.; Bangert, S.; Wiedelmann, submitted to the Protein Sequence Datal
                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein T18B22.110 - Arabidopsis thaliana (;Species: Arabidopsis thaliana (mouse-ear cress) C;Becies: Arabidopsis thaliana (mouse-ear cress) C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004 C;Accession: T47446
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A; Note: T18B22.110
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RESULT 10
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                                                                                GDTIDLIFNSEHQVLKNNTYW 114
                                                                                                                  WILPAIRNDLILLENQV------
                                                                                                                                               WKLEILQNGLYLIYGQVAPNANYNDVAPFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHV
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28.3%;
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Pred. No. 22;
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Pred. No. 23;
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hypothetical protein PAB2235 - Pyrococcus abyssi (strain Orsay) C;Species: Pyrococcus abyssi (c;Species: Pyrococcus abyssi C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change C;Accession: D75207

#text_change

09-Jul-2004

submitted to the EMBL Data Library, July A,Description: Pyrococcus abyssi genome &

sequence: 1999

insights

into

archaeal chromosome

str

R; anonymous,

Genoscope

D75207

A; Reference number: A75001 A; Accession: D75207 A; Description: Pyrococcus

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A;Cross-references: UNIPROT:Q54503; A;Experimental source: strains 5296
                                                                                                                 R;Dowson, C.G.; Hutchison, A.; Woodford, N.; Johnson, A.P.; George, R.C.; Proc. Natl. Acad. Sci. U.S.A. 87, 5888-5862, 1990
A;Title: Penicillin-resistant viridans streptococci have obtained altered A;Reference number: A35965; MUID:90332682; PMID:2377622
                                                                                                                                                                                                                                                                            RESULT 12
F35965
                                                           A;Status: preliminary; not compared with conceptual translation A;Molecule type: DNA
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A;Gene: PAB2235
A;Gene: PAB2235
C;Superfamily: Pyrococcus horikoshii hypothetical protein PH0196
                                             A;Residues:
                                                                                                                                                                                            penicillin-binding protein 2B - Streptococcus oralis (fragment)
C;Species: Streptococcus oralis
C;Date: 09-Nov-1990 #sequence_revision 09-Nov-1990 #text_change 09-Jul-2004
C;Accession: F35965
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C;Superfamily: penicillin-binding protein 2B
C;Keywords: antibiotic resistance; penicilli:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Coffey, T.J.; Dowson, C.G.; Daniels, M.; Spratt, B.G. submitted to the EMBL Data Library, March 1993
A;Description: Horizontal spread of an altered penicillin-binding
                                                                                                   A; Accession: F35965
                                                                                                                                                                                                                                                                                                                                                                                                                                                            δÃ
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A; Residues: 1-477 < COF>
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A;Accession: S32223
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A;Residues: 1-393 <KAW>
A;Cross-references: UNIPROT:Q9V291; GB:AJ248283; GB:AL096836; NID:g5457433; PIDN:CAB491
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Best Local
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Pred. No.
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Pred. No.
                   GB:M32226;
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A;Experimental source:
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A;Reference number: A43458; MUID:92407024; PMID:1527163 A;Accession: A43458
                                                                                                          replication protein A1 - African clawed frog
N;Alternate names: replication protein A 70K chain
C;Species: Kenopus laevis (African clawed frog)
C;Date: 10-Jun-1993 #sequence_revision 27-Oct-1995 #text_change
C;Accession: A43458; S24638
R;Adachi, Y; Laemmli, U.K.
J. Cell Biol. 119, 1-15, 1992
J. Cell Biol. 119, 1-15, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nature 390, 580-586, 1997
Nature 390, 580-586, 1997
A;Authors: Smith, H.O.; Venter, J.C.
A;Title: Genomic sequence of a Lyme disease spirochaete,
A;Title: Genomic sequence of a Lyme disease spirochaete,
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                                    A; Molecule type: mRNA; protein
                                                                                                                                                                                                                                               A43458
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   A;Cross-references:
                     A;Residues: 1-609 <ADA>
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 UNIPROT: Q01588;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.7%;
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Pred. No.
EMBL:X67240; NID:g65065;
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A;Molecule type: DNA
A;Residues: 1-1123 <ARN>
A;Residues: 1-1123 <ARN>
A;Cross-references: UNIPROT:Q9X083; GB:AE001760; GB:AE000512; NID:g4981510; PIDN:AAD3606
A;Experimental source: strain MSB8
                                                                                                                                                                                                                                                                                                                                                 A;Gene: TM0988
C;Superfamily: Pyrococcus abyssi hypothetical protein PAB0589
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 conserved hypothetical protein - Thermotoga maritima (strain MSB8)
c;Species: Thermotoga maritima
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: A72311
R;Nelson, K.E.; Clayton, R.A.; Gill, S.R.; Gwinn, M.L.; Dodson, R.J.; Haft, D.H.; Hid Garrett, M.M.; Stewart, A.M.; Cotton, M.D.; Pratt, M.S.; Phillips, C.A.; Richardson, C.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Note: sequence extracted from NCBI backbone (NCBIN:113390, NCBIP:113391)
A;Note: parts of the sequence determined by protein sequencing
C;Complex: Replication protein A is a trimer of 70K (A1), 32K (A2), and 14K (A3) chains.
C;Function:
A;Description: thought to be the eukaryotic equivalent of prokaryotic single-stranded DN A synthesis
A;Pathway: DNA replication initiation
C;Superfamily: replication protein A1
C;Keywords: DNA replication initiation; single-stranded DNA binding; trimer; zinc finger
F;472-494/Region: zinc finger CCCC motif
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A;Title: Evidence for lateral gene transfer between Archaea and Bacteria from genome A:Reference number: A72200; MUID:99287316; PMID:10360571
A;Accession: A72311
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25.7%; Pred. No. 94;
tive 21; Mismatches
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AATZ8927
QGTB42
QGMCG4
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QGIMB9
QSLFIL1
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CAE12167
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TNF5_CHICK
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FUNCTION: interacti and may m	SEQUENCE OF 9-177 SEQUENCE OF 9-177 TISSUE=Brain; MEDLINE=99156876; KWON B., Yu KY., Liu D., Wang SX. "Identification of necrosis factor re	SEQUENCE FROM NA MEDLINE=2288729 Clark H.F., Gur Chen J., Chow E Baton D., Foste Baton E., Liao Seshagiri S., S Vandlen R., Wath Yi S., Yu G., Y Godowski P., Gr "The secreted p effort to ident bioinformatics Genome Res. 13:	SEQUENCE FROM N.A. SEQUENCE FROM N.A. TISSUE=Umbilical vein MEDLINE=99175482; Pub Gurney A.L., Marsters Baldwin D.T., Gray A. Goddard A.D., Wood W. "Identification of a and its receptor, a h Curr. Biol. 9:215-218 [2]	HUMAN 12; 095 17-2001 17-2001 17-2004 17-2004 17-2004 17-2004 17-2004 11-2004 11-2004 11-2004 11-2004 11-2004 11-2004 11-2004	75 75 75 75	75.5 75.5 75.5 75.5 75.5 75.5
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274:6056-6 Cytokine thens between dulate T lydulate T l	0 7 15 15	KOM N.A. 897296; PubMed=1297531 897296; A.L., Abaya I how B., Chni C., Crow. Foster J., Grimaldi C. Foster J., Klimowski L. Liao D., Mark M., Robi S., Simmons L., Singh Watanabe C., Wieand G., Yuan J., Zhang M., Gray A.; ted protein discovery identify novel human tics assessment."; 13.2265-2270(2003).	A. vein; ly PubMed=1007; sters S.A., F. sters S.A., Down ay A.M., Down of a new ment of a new ment c, a human ort c, a human ort c, a human ort	ANDARD; 40, Createc 40, Last se 44, Last an ctor ligand ed ligand) UNQ149/PRO1; onyms=AITRL, an). Chordata a; Primates	4 3 3 5 5 2 3 2 5 9 2 3 2 2 9	390 3063 270 270 270 314 318 3890 3890
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-6051(1999). that binds to TNFRSF18/AITR/GITR. I n activated T lymphocytes and endot lymphocyte survival in peripheral t (Poftential).	7686; GL., Jang IK., Kim tivation-inducible prot	Baker K., Baldwi E., Baker K., Baldwi Ley C., Currell B., Gu Q., Hass P.E., Jin Y., Johnson S D., Smith V., Stins D., Woods K., Xie b Zhang Z., Goddard initiative (SPDI), secreted and transme	4428; Huang A., Pitti R.M., P., Brush J., Helde Rr K.P., Godowski P.J Per of the tumor necr	PRT; 177 AA. a) guence update) motation update) superfamily member 18 (MGITRL) (Activation-in 75) (GITRL, TL6; (GITRL, TL6; (Craniata; Vertebrata; (Catarrhini; Hominidae	VTS	Q615S8 Q8QXN2 Q72T60 Q8F293 AAS69768 Q31417 Q99U53 Q7A5M1 TWFA_FELCA
ITR. Important for endothelial cells eral tissues.	YJ., Xing L., ein of the tumor	in D., Brush J., Deuel B., Dowd P., Heldens S., S., Lee J., S., Lee J., Son J., Vagts A., MH., Yansura D., A., Wood W.I., a large-scale embrane proteins: a	Mark M., ns S., Schow P., , Ashkenazi A.; osis factor family	(Glucocorticoid- ducible TNF-related Euteleostomi;	Aar05186 unculture Aar05186 nanoarchaeu Aar39283 nanoarcha Q7wze8 shewanella	Q615s8 streptococc Q8qxn2 sugarcane m Q72t60 leptospira Q8f293 leptospira Aas69788 leptospir Q31417 latimeria c Q39u53 staphylococ Q7a5m1 staphylococ Q7a5m1 staphylococ Q7af64 upcultured

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SUBUNIT: Homotrimer (Potential).

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RESULT 2
AAH69111
ID AAH6
AC AAH6
AC AAH6
DT 24-w
DT 24-w
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DT TMFS
GN TNFS
GN TNFS
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Matches 126
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GO; GO:0006916; P:anti-apoptosis; TAS.
GO; GO:0007267; P:cell-cell signaling; TAS.
GO; GO:0007165; P:signal transduction; TAS.
InterPro; IPR006052; TNF family.
InterPro; IPR008983; TNF like.
PROSITE; PS00251; TNF 1; FALSE NEG.
PROSITE; PS0049; TNF 2; FALSE NEG.
Cytokine; Glycoprotein; Signal-anchor; Transmu
   TISSUE=Synthetic constructs;

MEDLINE=22388257; PubMed=12477932;

Strausberg R.L., Feingold E.A., Grouse L.H.,

Klausner R.D., Collins F.S., Wagner L., Shenn
Altschul S.F., Zeeberg B., Buetow K.H., Schae
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24-MAY-2004 (TrEMBLrel. 27,
24-MAY-2004 (TrEMBLrel. 27,
24-MAY-2004 (TrEMBLrel. 27,
TNFSF18 protein (Fragment).
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Mammalia; Eutheria;
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CARBOHYD
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
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EMBL; AV358868; AAQ89277.1; -.
EMBL; AF117713; AAD19695.1; -.
Genew, HGNC:11932; TNFSF18.
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INDUCTION: Up-regulated after stimulation by lipopolysaccharides.
SIMIDARITY: Belongs to the tumor necrosis factor family.
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                                                                                                                                                                                                                                                                                                                                                                            121
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Extracellular (Potential).
N-linked (GlcNAc. . .) (Potential)
N-linked (GlcNAc. . .) (Potential)
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Signal-anchor for type II membrane
                                                                                                                          Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 675; DB 1;
Pred. No. 1.1e-57;
                                                                                                                                          Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3D78CE6B90F4C9E3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                   177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (See http://www.isb-sib.ch/announce/
L.H., Derge J.G.,
Shenmen C.M., Schuler
Schaefer C.F., Bhat N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Transmembrane.
                                                                                                                                                                                                                                                                   A
                                                                                                                                                                                                      update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 177;
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RESULT
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X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Scheetz T.E.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Scheetz T.E.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Scheetz T.E.,

X Hopkins R.F., Jordan H., Moore T., Abramson R.D., Mullahy S.J.,

X Haha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

X Robards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

X Hilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

X Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

X Hilling M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2004 (TrEMBLrel.
01-JUN-2004 (TrEMBLrel.
01-JUN-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; BO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Scheetz T.E., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.M., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S., Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.E.,
                                                                                                                                                                                                                                                                               TISSUE=PCR rescued clones;
                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH69319;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones S.J., Marra M.A.;
"Generation and initial analysis
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strausberg R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Synthetic constructs;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NPQFIS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILLA
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                                                                                                                                                                                                                                                                                                                                                                                                 (Human).
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                                                                                                                                                                                                                                                                                                                                                           Chordata;
Primates;
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27,
27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 675; DB 2;
Pred. No. 1.1e-57;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
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QRESULT QROYGEZ ID YGZ 
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Best Local S
Matches 126
Query Match
Best Local Similarity
                                                                        costimulatory for T cells.";

proc. Natl. Acad. Sci. U.S.A. 100:15059-15
EMBL; AV234223; AAA089011.1; -.
EMBL; AJ577579; CAE12166.1; -.
EMBL; AJ577580; CAE12166.1; -.
MGD; MG1:2673064; Thisfils
GO; GO:0016020; C:membrane; IEA.
GO; GO:00069164; F:tumor necrosis factor re
GO; GO:00069165; P:immune response; IEA.
InterPro; IPR008983; TNF like.
InterPro; IPR008983; TNF subf;
ProDom, PD002012; TN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blakesley R.W., Touchman J.W., Gr
Rodriguez A.C., Grimwood J., Schm
Krzywinski M.I., Skalska U., Smai
Jones S.J., Marra M.A.;
"Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name=Tnfsf18; Synonyms=tnfsf
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GITR ligand
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01-JUN-2003 (TrEMBLrel.
01-JUN-2003 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q80YG2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bianchini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                               Waldmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tone M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=DBA/2J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                  "Mouse glucocorticoid-induced tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4
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BC069319; AAH69319.1; -.
NCB 177 AA; 20307 MW; 3D78CE6B90F4C9E3 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FROM N.A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tone Y., Adams
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nocentini
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Rodentia;
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56.7%;
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EMBL/GenBank/DDBJ databases.
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Sciurognathi; Muridae;
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No. 1.1e-57;
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                                                                           CAE12167;
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                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                     "Mouse glucocorticoid-induced tumor necrosis factor costimulatory for T cells."; proc. Natl. Acad. Sci. U.S.A. 100:15059-15064(2003) EMBL; AJ577579; CAE12166.1; -
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                                                                                                                                                                                                                                                                                                      Alternative splicing.
SEQUENCE 173 AA; 19745 MW;
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                                Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                     GITR ligand.
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                       NCBI_TaxID=10090;
                                                             TNFSF 18
         SEQUENCE FROM N.A.
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Rodentia;
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Rodentia;
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4; Mismatches 39
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Sciurognathi;
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Sciurognathi; Muridae;
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01-OCT-2003
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01-OCT-2003
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EMBL; AY267900; AAP96745.1; ---
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PubMed=14647196;
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167
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                                                                      APFVVQIYKKNDVLQTLMNDFQILPIGGVYELHAGDNIYLKFNSKDHIQKNNTYWGIILM
                                                                                                         APFEVRLYKNKDMIQTLINKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILL
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                                                                                                                                                     TVIESCMVKFELSSSKWHMTSPKPHCVNTTSDGKLKILQSGTYLIYGQVIPVDKKYIKDN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e splicing.
173 AA; 19745 MW;
                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Adams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chordata; Craniata; Vertebrata; Buteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                   55.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              56.7%;
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                                                                                                                                                                                                                                               14;
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                                                                                                                                                                                                                                         Score 344; DB 2; ]
Pred. No. 1.8e-25;
4; Mismatches 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 348; DB
Pred. No. 7.5e
14; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Frewin M.R.,
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                                                                                                                                                                                                                                                                                   Length 173;
                                                                                                                                                                                                                                                                                                                                 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                              binding;
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                                                                                                                                                                                                                                             Indels
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RESULT Q7TS55

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Best Local
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                                                                                                                                                              02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                          AAQ55265;
AAQ55265;
                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                         02-MAR-2004 (TrEMBLrel. 27, I 02-MAR-2004 (TrEMBLrel. 27, I Glucocorticoid-induced-tumor
SEQUENCE FROM N.A
                                    NCBI_TaxID=10090;
                                                                                                  Mus musculus
                                                                                                                            GITRL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Receptor.
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yu K.-Y., Min S.S., Youn B.-S.;
Submitted (AUG-2003) to the EMBL/GenB
EMBL; AY320040; AAP70494.1; -
EMBL; AY359852; AAQ55265.1; -
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004872; F:receptor activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-2003
05-JUL-2004
GITR ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q7TS55;
Q7TS55;
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=22883945; PubMed=14521928;
Yu K.Y., Kim H.S., Song S.Y., Min S.S., Jeong J.J., Youn B.S.;
"Identification of a ligand for glucocorticoid-induced tumor not
factor receptor constitutively expressed in dendritic cells.";
Biochem. Biophys. Res. Commun. 310:433-438(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=129/J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ProDom;
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Ji H., Terhorst C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=Gitrl;
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GO:0006955; P:immune response;
GPPO; IPR003636; INF subf.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PD002012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APFEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PDLPFIS
                                                                                                                                                                                                                                                                                                                                                                                           ANPOFIS
                                                                                                                                                                                                                                                                                                                                                                                                                           APFVVQIYKKNDVLQTLMNDFQ1LPIGGVYELHAGDNIYLKFNSKDHIQKTNTYWGIILM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (JUN-2003) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               173
                                                                                                                                                                                                     (TrEMBLrel. 27,
                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                    (Mouse).
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(TrEMBLrel. 25, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
(Glucoccorticoid-induced-tumor necrosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AΑ;
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                                                         Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19732 MW;
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Rodentia;
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                                                                                                                                                          Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 342; DB 2;
Pred. No. 2.9e-25;
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                                                                                                                                           necrosis
                                                 Craniata; Vert
Sciurognathi;
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Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0F08494CACF424D2
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                                                                                                                                         factor receptor
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                                                                             Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor
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thi; Muridae;
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                                                     Muridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             binding; IEA.
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                                                                             Euteleostomi;
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; Murinae; Mus
                                                       Murinae; Mus
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TRESULT 10
TNP5
AC Q918E
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OS Galli
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Best Local S
Matches 71
                                                                                                                                                                                                                                                                                                                                                                         STRAIL-white leghorn; TISSUE-Spleen;

A Tregaskes C.A., Young J.R., Burnside J.;

The "Cloning of a putative chicken CD40 ligand.";

"Cloning of a putative chicken CD40 ligand.";

Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.

- FUNCTION: Cytokine that binds to TNFRSF5. Mediates B-cell

- FUNCTION: Cytokine that binds to TNFRSF5. Mediates B-cell

- FUNCTION: Cytokine that binds to TNFRSF5. Mediates B-cell

- FUNCTION: Cytokine that binds to TNFRSF5. Mediates B-cell

- Collass switching (By similarity).

- IssueUNIT: Homotrimer (By similarity).

- IssueUNIT: Homotrimer (By similarity).

- IssueUNIT: The soluble form derives from the membrane form by proteolytic processing (By similarity).

- IssueUNIT: The soluble form derives from the membrane form by proteolytic processing (By similarity).
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Submitted
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Yu K. Y., Kim H.S., Song S.Y., Min S.S., Jeong J.J., Youn B.S.;
"Identification of a ligand for glucocorticoid-induced tumor ne
factor receptor constitutively expressed in dendritic cells.";
Biochem. Biophys. Res. Commun. 310:433-438(2003).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2003 (Rel. 41, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Tumor necrosis factor ligand superfamily member
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Q918D8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=TNFSF5; Synonyms=CD40LG, Gallus gallus (Chicken).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L) (CD154 protein).
                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commentities requires a license agreement (See http://www.isb-sib.ch/an
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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    EMBL; AJ243435;
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                                                                                an email to license@isb-sib.ch).
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(AUG-2003) to the
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Pred. No. 2.9e-
14; Mismatches
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                                                                                                                            (See http://www.isb-sib.ch/announce/
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GO; GO:0016021; C:integral to membrane; ISS.
GO; GO:0005174; F:CD40 receptor binding; ISS.
GO; GO:0005174; F:CD40 receptor binding; ISS.
GO; GO:00042100; P:B-cell proliferation; ISS.
GO; GO:0006954; P:inflammatory response; ISS.
GO; GO:0007159; P:leukocyte cell adhesion; ISS.
GO; GO:0007159; P:platelet activation; ISS.
R InterPro; IPR003263; TNF_5.
R InterPro; IPR008983; TNF_1ike.
R InterPro; IPR008983; TNF_1ike.
R InterPro; IPR008983; TNF_Subf.
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PRINTS; PR01702; CD40LIGAND.
ProDom; PD00860; TNF 5; 1.

ProDom; PD002012; TNF 5; 1.

PR0SITE; PS000251; TNF 2; 1.

PR0SITE; PS50049; TNF 2; 1.

Cytokine; Glycoprotein; Signal.
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                                                           Gloeckner G., Eichinger L., Szafranski K., Pachebat J., Dear P., Lehmann R., Baumgart C., Parra G., April J.F., Guigo R., Kumpf K., Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noegel A.A.; "Sequence and analysis of chromosome 2 of Dictyostelium discoideum."; Nature 418:79-85(2002).
                                                                                                                                                                                                                                                                                 Dictyostelium discoideum (Slime mold)
Eukaryota; Mycetozoa; Dictyosteliida;
                                                                                                                                                                                                                                                                                                                            Hypothetical protein.
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01-MAR-2004 (TrEMBLrel.
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STRAIN=AX4;
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SEQUENCE FROM N.A
STRAIN=AX4;
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Gloeckner G., Eichinger L., Szafra
Lehmann R., Baumgart C., Parra G.,
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member 5, soluble form (By similarity).
Cytoplasmic (Potential).
Signal-anchor for type II membrane
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Extracellular (Potential).
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annotation update)
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RESULT 13
Q8IAW5
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Best Local Similarity
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Matches 27
           Q8IAW5;
Q8IAW5;
01-MAR-2003
01-MAR-2003
01-MAR-2003
  Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q896D6 PRELIMINARY; PRT; 270 AA.
Q896D6;
Q1-JUN-2003 (TrEMBLrel. 24, Created)
O1-JUN-2003 (TrEMBLrel. 24, Last sequence update)
O1-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Kila protein, putative phage-related DNA binding protein.
Name=kila, OrderedLocusNames=CTC01071;
                                                                                                                                                                                                                                                                                                                                 Complete proteome. SEQUENCE 270 AA;
                                                                                                                                                                                                                                                                                                                                                      Proc. Natl. Acad. Sci. U.S.A. 100:1316-1321 (2003)
EMBL; AE015939; AA035654.1; -.
InterPro; IPR003497; BRO_N.
Pfam; PP02498; Bro-N; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=22457253; PubMed=12552129;
Brueggemann H., Baeumer S., Fricke W.F., Wiezer A., 1
Decker I., Herzberg C., Martinez-Arias R., Merkl R.,
                                                                                                                                                                                                                                                                                                                                                                                                        "The genome sequence of Clostridium tetani, the causative tetanus disease.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=Massachusetts / E88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Clostridium tetani.
Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Gottschalk G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Baumgart C.;
Submitted (FEB-2004) to the EMEMBL; ACI16551; AAO52204.1; -.
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=1513;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clostridium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29 NKVSDWKLEILQNGLYLIYGQVAENANYNDYAPFEYRLYKNKDMIQTLINKSKIQNVGGT
                                                                                                                              NA
                                                                                                                                                       A
                                                                                                                                                                         LNKLLHELKVQYKQNDQWLLYKEHSGKGYTHSETIDIVRSDGRPDVKVNYKWTQKGRLFL
                                                                                                                                                                                                                         KAKNKMLELQNKQKEQIIĞELKPRADYTD-----RILKNKGLV-TITQIAKDYGMTĞİG 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                  KASDMKTEITÖNGTATTÄGÖAMANANANABAEAKTAKNKDWIÕLTINKSKIÖNAGGI-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YELHVGDTIDLIFNSEHQVLKNNTYWGIILLAN 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NETEDSKLTF----IXIIY-----NIEKQDVQPQTVIEFKTDEINEMITHKFQIENNDDS
      (TrEMBLrel. 23, Created)
(TrEMBLrel. 23, Last sequence update)
(TrEMBLrel. 23, Last annotation update)
                                                                                                                              248
                                                                                                                                                       121
                                                                                                                                                                                                                                                                             12.7%; Score 85.5; I
ilarity 25.4%; Pred. No. 4.7;
Conservative 18; Mismatches
                                                               PRELIMINARY;
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 protein
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                                                                                                                                                                                                                                                                                                                                 31533 MW;
  PF08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.8%; Score 86.5; 29.0%; Pred. No. 7
_0083.
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                                                                                                                                                                                                YELHVG-----DTIDLIFNSEHQVLKNNTYW---GIILL 119
                                                                                                                                                                                                                                                                                                                              C378AB524C0133A8 CRC64;
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                                                             599
                                                                                                                                                                                                                                                                                            4.7;
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RESULT 14
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                                                                                                                                                                                                                              R MEROPS; C06.001; -..

R G0; G0:0019028; C:viral capsid; IEA.

R G0; G0:0005524; F:ATP binding; IEA.

R G0; G0:0005524; F:ATP binding; IEA.

R G0; G0:0008126; F:ATP-dependent helicase activity; IEA.

R G0; G0:0008127; F:cysteine-type endopeptidase activity; IEA.

G0; G0:0004197; F:hydrolase activity; IEA.

R G0; G0:000373; F:RNA binding; IEA.

R G0; G0:000373; F:RNA binding; IEA.

R G0; G0:0003768; F:RNA-directed RNA polymerase activity; IEA.

R G0; G0:00035198; F:structural molecule activity; IEA.

R G0; G0:00055198; F:structural molecule activity; IEA.

R G0; G0:0005508; P:proteolysis and peptidolysis; IEA.

R G0; G0:0006508; P:proteolysis and peptidolysis; IEA.

R G0; G0:0006508; P:viral genome replication; IEA.

R G0; G0:0006509; P:viral genome replication; IEA.

R G0; G0:0019079; P:viral genome replication; IEA.
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Best Local
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Characterisation of potyviruses from sugarcane and maize in China.";
"Characterisation of potyviruses from sugarcane and maize in China.";
"Arch. Virol. 147:1237-1246(2002).
-!- SIMILARITY: Belongs to the potyviruses polyprotein family.
                                                       InterPro;
                                                                                      InterPro;
                                                                                                               InterPro;
                                                                                                                                        InterPro;
                                                                                                                                                                    InterPro;
                                                                                                                                                                                                InterPro;
                                                                                                                                                                                                                         InterPro;
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MEDLINE=22105130; PubMed=12111432;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Viruses; ssRNA positive-strand viruses, no DNA stage; Potyviridae;
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01-JUN-2002 (TrEMBLrel. 21,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sugarcane mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plasmodium falciparum (isolate 3D7).
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
NCBI_TaxID=36329;
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                           prolis45; DEAD/DEAH N.
FIR001650; Helicase C.
FIR001330; Peptidase C4.
FIR001456; Peptidase C6.
FIR002540; Pept_S30 Poty_F
FIR002540; Pept_S30 Poty_F
FIR001592; Poty_Coat.
FIR007095; RNA_pol_DS_PS.
FIR001205; RNA_pol_DS_PS.
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RNA_pol_DS_PS.
RNA_pol_P3D.
RNA_pol_PSvir.
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22.9%; Pre
7ative 24;
                                                                                                    Peptidase_C6.
Pept_S30_Poty_P1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 83; DB 2
Pred. No. 21;
24; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Last sequence update)
Last annotation updat
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Best Local S
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SMART; SMO049
SMART; SMO049
ATP-binding;
CHAIN 10
CHAIN 11
CHAIN 17
CHAIN 17
CHAIN 17
CHAIN 19
CHAIN 22
CHAIN 27
CHAIN 22
CHAIN 27
InterPro;
InterPro;
Pfam; PF00
                                                                            InterPro;
InterPro;
InterPro;
                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Dihydrolipoamide dehydrogenase (EC 1.8.1.4).

Name-acoL; ORFNames=E79727 2536;
Bacillus thuringiensis serovar konkukian str. 97-27
Bacteria; Firmicutes; Bacillales, Bacillaceae; Bacillus thuringiensis serovar konkukian.
                                                                                                                                                                                                                     EMBL;
                                                                                                                                                               InterPro;
                                                                                                                                                                                         InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=281309;
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Pfam;
Pfam;
Pfam;
Pfam;
Pfam;
Pfam;
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                 oxidoreductase family.

MBL, AE017355; AAT61350.1; --
nterPro; IPR001327; FAD_Dyr_redox.

nterPro; IPR000815; Hg_reductase.

nterPro; IPR0006258; Lipoamide_dh.

nterPro; IPR000103; PyrIdine_redox_1

nterPro; IPR000100; Pyr_redox_1

nterPro; IPR001100; Pyr_redox_dim.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
                                                                                                                                                                                                                                                                   SIMILARITY: Belongs to the class-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; PF00270; DEAD; 1.
1; PF00271; Helicase C; 1.
1; PF00863; Peptidase C6; 1.
1; PF00851; Peptidase C6; 1.
1; PF00767; Poty Coat; 1.
1; PF01577; Poty Pl; 1.
1n; PF00680; RNA dep RNA pol; 1
NTS; PR00966; NIAPOTYPTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             397
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SM00490; HELICc; 1
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  redox;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         693
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NIa-VPg protein.
NIa-protease protein.
Pl protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 83; DB 2;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              coat protein.
HC-Pro protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NIb polymerase protein
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6K1 protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E6019CBD1E80B59D CRC64;
                                                                                                                                                                                                                                                                                                                         is a redox-active disulfide bond
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                                                                                                                                                                                                                                                                      nucleotide-disulfide
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s., Okinaka
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Okinaka R.,
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Best Local (
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PRINTS; PR00368; FADDARR.
PRINTS; PR00945; HGRDTASE.
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PRINTS; PR00469; PNDEDTASEII.
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256
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                                                                                                                  Similarity
ANPEYV
                ANPOFI 125
                                    AP-
                                                 APPEVRLYKNKDMIQTLTNKSKIQNVGGTYELHVGDTIDLIFNSEHQVLKNNTYWGIILL
                                                                      EPAELPFAPFDGKWILNSSHAMSLENIPK-SLLIVGGGVIGCEFASIYSRLGTKVTIVEM
                                                                                        EPCMAKFGPLPSKWQMASSEPPCVNKVSDWKLEILQNGLY-----LIYGQVAPNANYNDV
                                  -QLLPGEDKDIAQIL--KEKLESDG--VEIFTGAALKGLNNYKKQA--SFEYKGSIQE
                                                                                                           Conservative
261
                                                                                                                  11.8%;
                                                                                                          25;
                                                                                                                  Score 79.5;
Pred. No. 3
                                                                                                          Mismatches
                                                                                                                  33;
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119 255

59 202

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Search completed: November 2, 2004, 12:58:45
Job time: 194 secs